

**Exploring Image and Video by Classification and Clustering on  
Global and Local Visual Features**

By

Le Lu

M.S.E. Johns Hopkins University 2004

B.E. Beijing Polytechnic University 1996

A Dissertation submitted to  
The Faculty of  
Whiting School of Engineering  
of The Johns Hopkins University in partial satisfaction  
of the requirements for the degree of Doctor of Philosophy

Fall 2006

Dissertation directed by

Gregory D. Hager

Professor of Computer Science



# Abstract of Dissertation

Images and Videos are complex 2-dimensional spatial correlated data pattern or 3-dimensional spatial-temporal correlated data volumes. Associating these correlated relationships of visual data signals (acquired by imaging sensors) with high-level semantic human knowledge is the core challenging problem of pattern recognition and computer vision. Finding the inter-correlated relationships across multiple different images or videos, given a large amount of data in similar or non-similar scenarios, without direct annotation from human concepts, is another self-organized data structuring issue. From the previous literature and our own research work [9, 11, 12, 13, 14, 15], via computing machines as tools, there are a lot of efforts trying to address these two tasks statistically, by making good use of recently developed supervised (a.k.a. Classification) and Unsupervised (a.k.a. Clustering) statistical machine learning paradigms [1, 5, 26, 7, 165, 19, 145, 2, 18].

In this dissertation, we are particularly interested to study four specific yet important computer vision problems involving partitioning, discriminative multiple-class classification and online adaptive appearance learning, depending on statistical machine learning techniques. Our four tasks are based on extracted global and local visual appearance patterns from both image and video domains respectively. First, we develop new unsupervised clustering algorithm to partition temporal video structure (a.k.a. video shot segmentation) [12]. Second, we detect and recognize spatial-temporal video volumes as action unites through trained 3D surface model and multi-scale temporal searching [60]. The 3D surface based action model is obtained as output of the learning process of texton-like [121] intermediate visual representations, using sequentially adapted clustering from figure-segmented image sequences [9]. Third, we train discriminative multi-modal probabilistic density classifiers to detect semantic material classes from home photo in a soft

classification manner. Learning photo categories is then based the global image features extracted from the material class-specific density response maps by random weak classifier combination [68, 103, 162] to handle the complex photo-feature distribution in high dimensions [11]. Fourth, we propose a unified approach for close-field (medium to high resolution) segmentation based object matching and tracking (a.k.a video matting [73, 124, 161]), and far-field localization based object tracking [?, 96]. The main novelty exists on two folds: our framework that allows very flexible image density matching function construction [5, 165, 69, 100, 106, 11, 87, 86, 71]; our bi-directional appearance consistency check algorithm which has been demonstrated to maintain an effective object-level appearance models under different severe changing, occluding and deformable appearance situations and camera imaging conditions, but only require very simple nonparametric similarity computations [14].

# Acknowledgments

Thank some people here.

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# Introduction

## 0.1 Motivation

Using *feature based* (ie. interest image features that satisfy some metrics of geometric or photometric invariance [99, 110, 126, 131, 130]) or *direct image methods* (ie. derivatives or differences of image intensity patterns [62, 97, 109]) for 3D object/scene model construction [108, 154], image content retrieval [142, 75], object recognition [83, 126], video structure matching/parsing [140, 150, 149] and automatic mosaics generation [70] have been well exploited in computer vision community during the last decade. The advantage of *feature based methods* is that some geometric and photometric invariance can be encoded into the feature design and detection process. Thus features can be repeatedly detected in a relatively more stable manner with respect to image changes, illumination variations and random noises, than *direct image methods*. Additionally, *feature based methods* is usually more robust with occlusions, due to its local part-based representation. On the contrary, *direct image methods* will prevail when image features are hard to find or the predefined feature detection principles are not coincident with the given vision task. Without extra efforts on designing and finding features, *direct image methods* can be performed very fast and often in realtime [97, 109]. As a summary, *feature based methods* are more likely to be employed by representing high-resolution visual scenes with many distinct "corner" like local features; while *direct image methods* have more privileges by characterizing low-resolution imagery, textureless or homogenous regions, highly repeated textures and images containing dominant "edge or ridge" like features<sup>1</sup>.

In this proposal, we represent images using sets of regularly or irregularly spatially sampled rectangle

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<sup>1</sup>Edge or ridge features can be conveniently computed using simple image gradient operators [62, 97] or filter banks [128, 129]

subregions of interest (ROI), ie. image patches, as an intermediate solution between *feature based methods* and *direct image method*. The image patches have much lower dimensionality than a regular sized image which makes the statistical learning problem much easier. The pool of patches can be drawn randomly from larger labelled image regions, as many as what we need. Sufficient large sets of training image patches are guaranteed. Any given image can be modelled as a distribution of its sampled image patches in the feature space, which is much more flexible than direct method of modelling the image itself globally.

We demonstrate its representative validity by classifying a large photo database with very diverse visual contents [11] into scene categories and segmenting nonrigid dynamic foreground/background regions in video sequences [?] with satisfying results. More precisely, we build a probabilistic discriminative model for scene recognition which is learned over thousands of labelled image patches. The trained classifier performs the photo categorization task very effectively and efficiently [11]. Breaking images into a chuck of patches enables us to build a flexible, conceptually simple and computationally efficient discriminative classifier with good generalization comprehensive modelling capacity. Our recognition rate [11] is one of the best reported results<sup>2</sup> [127, 144, 66, 151, 81].

The challenging computer vision task of video foreground/background segmentation under dynamic scenes further validates our concept of image patch based representation. Our method generates good results on several difficult dynamic close-view video sequences captured with a moving camera, while other state-of-art algorithms [146, 133, 161, 124] mainly work on static or quasi-dynamic scenes. In our approach, distributed foreground/background image regions of very complex visual appearances are statistical-sufficiently sampled and formed into two nonparametric foreground/background appearance models. Many popular statistical clustering, density estimation techniques and dimension reduction algorithms [5, 165, 84, 87, 69, 106] can be employed to build the appearance models. A simple heuristic is also proposed in [?] on how to extract patches adaptively from any given image according to its spatial distribution of visual content complexity, by leveraging a general image segmentor [82]<sup>3</sup>. This spatial-sampling adaptiv-

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<sup>2</sup>Because there is no publicly available benchmark photo database for scene categorization, the above mentioned algorithms are tested with each individual photo database.

<sup>3</sup>Any image segmentor with reasonable performance can be used in our work separately or jointly.

ity is inspired by the idea that homogeneous image regions can be characterized by fewer image subregion samples, while irregularly textured image regions need more representative image patch samples. It condenses the size of the required representative data samples and decreases the algorithm's computational load as well. In the following, we provide details on the problem statement, algorithm description, preliminary experimental results and future extension plans for the two computer vision tasks mentioned above: *1, A Two-level Approach For Scene Recognition* and *2, Sampling, Partitioning and Aggregating Patches for Foreground/Background Labelling on Video Sequences.*

# Chapter 1

## Partitioning Temporal Video Structure: A Combined Central and Subspace Clustering Approach

*Content-based video exploration and understanding, as stated in [20], includes visual content analysis, video structure parsing, video summarization and indexing. In this chapter, we focus on providing a new machine learning solution for temporal video structure segmentation [6, 31, 32, 27, 17, 16, 3, 30] based on the temporal trajectory of frame-wise global visual features from principal component analysis (PCA) [5, 165].*

*On the other side, central and subspace clustering methods are at the core of many segmentation problems in computer vision. However, both methods fail to give the correct segmentation in many practical scenarios, e.g., when data points are close to the intersection of two subspaces or when two cluster centers in different subspaces are spatially close. In this paper, we address these challenges by considering the problem of clustering a set of points lying in a union of subspaces and distributed around multiple cluster centers inside each subspace. We propose a generalization of Kmeans and Ksubspaces that clusters the data by minimizing a cost function that combines both central and subspace distances. Experiments on synthetic data compare our algorithm favorably against four other clustering methods. We also test our algorithm on*

*computer vision problems such as face clustering with varying illumination and video shot segmentation of dynamic scenes.*

## 1.1 Introduction

### 1.1.1 Review on Clustering Methods

Many computer vision problems require the efficient and effective organization of huge-dimensional data for information retrieval purposes. Unsupervised learning, mostly clustering, provides a way to handle these challenges.

Central and subspace clustering are arguably the most studied clustering problems. In *central clustering*, data samples are assumed to be distributed around a collection of cluster centers, e.g., a mixture of Gaussians. This problem shows up in many vision tasks, e.g., image segmentation, and can be solved using techniques such as Kmeans [?] or Expectation Maximization (EM) [40].

In *subspace clustering*, data samples are assumed to be distributed in a collection of subspaces. This problem shows up in various vision applications, such as motion segmentation [24], face clustering with varying illumination [8], temporal video segmentation [25], etc. Subspace clustering can also be used to obtain a piecewise linear approximation of a manifold [29], as we will show in our real data experiments<sup>1</sup>. Existing subspace clustering methods include Ksubspaces [8] and Generalized Principal Component Analysis (GPCA) [25]. Such methods do not enforce a particular distribution of the data inside the subspaces. Methods such as Mixtures of Probabilistic PCA (MPPCA) [23] further assume that the distribution of the data inside each subspace is Gaussian and use EM to learn the parameters of the mixture model and the segmentation of the data.

Unfortunately, there are many cases in which neither central nor subspace clustering individually are appropriate. In motion segmentation, for example, there are two motion subspaces where each of them contains two moving objects and two objects from different motion subspaces are spatially close during

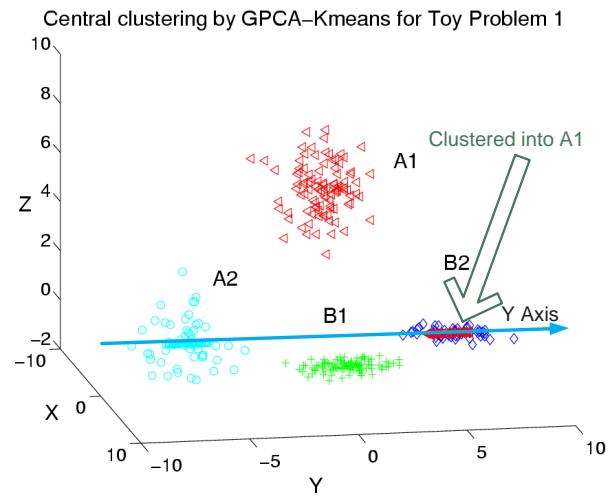
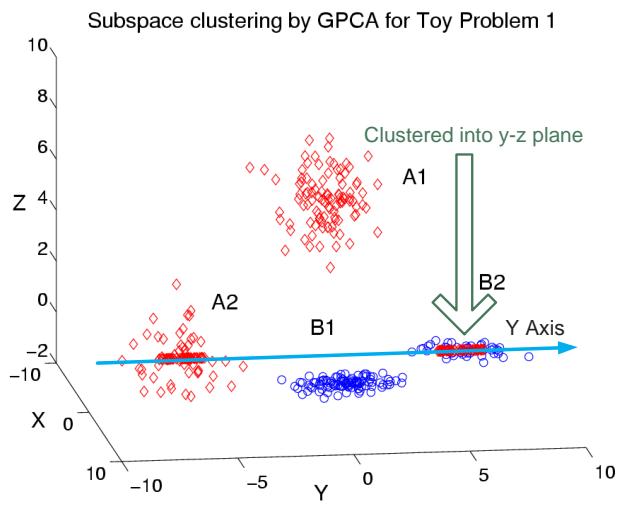
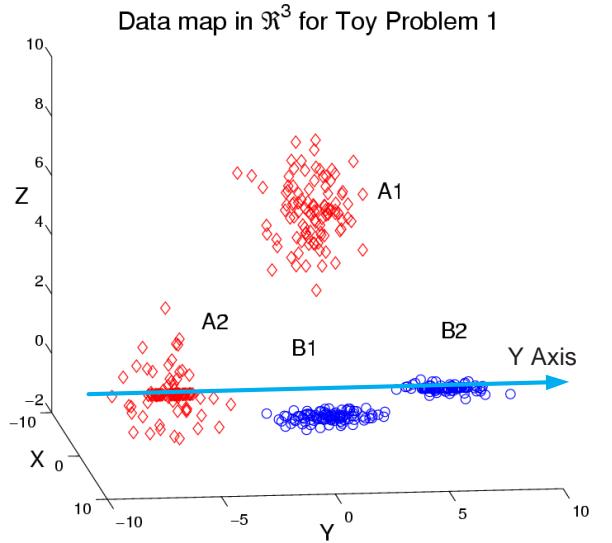
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<sup>1</sup>While techniques for learning manifolds from data already exist, e.g., [29], manifold parsing is a very difficult machine learning problem and has not been so well studied.

moving. If one is interested in grouping based on motion only, one may argue that the problem can be solved by subspace clustering of motion subspace alone. However, if one is interested in grouping the individual objects, the problem can not be well solved using central clustering of spatial locations alone, because the two spatially close objects under different subspaces can confuse central clustering. In this case, some kind of combination of central and subspace clustering should be considered.

### 1.1.2 Failure or Success: Two Toy Problems?

subspace clustering fails when the data set contains points close to the intersection of two subspaces, as shown by the example in Figure 1.1. Similarly, central clustering fails when two clusters in different subspaces are spatially close, as shown by the example in Figure 1.2.



**Figure 1.1:** **Top:** A set of points in  $\mathbb{R}^3$  drawn from 4 clusters labeled as  $A_1$ ,  $A_2$ ,  $B_1$ ,  $B_2$ . Clusters  $B_1$  and  $B_2$  lie in the x-y plane and clusters  $A_1$  and  $A_2$  lie in the y-z plane. Note that some points in  $A_2$  and  $B_2$  are drawn from the intersection of the two planes (y-axis). **Middle:** Subspace clustering by GPCA assigns all the points in the y-axis to the y-z plane, thus it misclassifies some points in  $B_2$ . **Bottom:** Subspace clustering using GPCA followed by central clustering inside each plane using Kmeans misclassifies some points in  $B_2$ .

In section 1.5, Through the visualization of real video data in their feature space, we find that data samples in  $\mathbb{R}^3$  are usually distributed on complex shaped curves or manifolds, as shown later, in Figure 1.7 and 1.8. With encouraging clustering results, our experiments also validate that subspace clustering can be effectively used to obtain a piecewise linear approximation of complex manifolds.

### 1.1.3 Proposed Clustering Method

In this paper, we propose a new clustering approach that combines both central and subspace clustering. We obtain an initial solution by grouping the data into multiple subspaces using GPCA and grouping the data inside each subspace using Kmeans. This initial solution is then refined by minimizing an objective function composed of both central and subspace distances. This combined optimization leads to improved performance of our method over four different clustering approaches in terms of both clustering error and estimation accuracy. Real examples on illumination-invariant face clustering and video shot detection are also performed. Our experiments also show that combined central/subspace clustering can be effectively used to obtain a piecewise linear approximation of complex manifolds.

## 1.2 Algorithm: Combined Central-Subspace Clustering

Let  $\{\mathbf{x}_i \in \mathbb{R}^D\}_{i=1}^P$  be a collection of  $P$  points lying approximately in  $n$  subspaces  $S_j = \{\mathbf{x} : B_j^\top \mathbf{x} = 0\}$  of dimension  $d_j$  with normal bases  $\{B_j \in \mathbb{R}^{(D-d_j) \times D}\}_{j=1}^n$ . Assume that within each subspace  $S_j$  the data points are distributed around  $m_j$  cluster centers  $\{\mu_{jk} \in \mathbb{R}^D\}_{j=1 \dots n}^{k=1 \dots m_j}$ . In this paper, we consider the following problem:

**Problem 1** (Combined central and subspace clustering). *Given  $\{\mathbf{x}_i\}_{i=1}^P$ , estimate  $\{B_j\}_{j=1}^n$  and  $\{\mu_{jk}\}_{j=1 \dots n}^{k=1 \dots m_j}$ .*

When  $n = 1$ , Problem 1 reduces to the standard central clustering problem. A popular central clustering method is the Kmeans algorithm, which solves for the cluster centers  $\mu_k$  and the membership of the  $i$ th point

to the  $k$ th cluster center  $w_{ik} \in \{0, 1\}$  by minimizing the within class variance

$$J_{KM} \doteq \sum_{i=1}^P \sum_{k=1}^{m_1} w_{ik} \|\mathbf{x}_i - \mu_k\|^2. \quad (1.1)$$

Given the cluster centers, the optimal solution for the memberships is to assign each point to the closest center. Given the memberships, the optimal solution for the cluster centers is given by the means of the points within each group. The Kmeans algorithm proceeds by alternating between these two steps until convergence to a local minimum.

When  $m_j = 1$  and  $n > 1$ , Problem 1 reduces to the classical subspace clustering problem. As shown in [8], this problem can be solved with an extension of Kmeans, called Ksubspaces, which solves for the subspace normal bases  $B_j$  and the membership of the  $i$ th point to the  $j$ th subspace  $w_{ij} \in \{0, 1\}$  by minimizing the cost function

$$J_{KS} \doteq \sum_{i=1}^P \sum_{j=1}^n w_{ij} \|B_j^\top \mathbf{x}_i\|^2 \quad (1.2)$$

subject to the constraints  $B_j^\top B_j = \mathcal{I}$ , for  $j = 1, \dots, n$ , where  $\mathcal{I}$  denotes the identity matrix. Given the normal bases, the optimal solution for the memberships is to assign each point to the closest subspace. Given the memberships, the optimal solution for the normal bases is obtained from the null space of the data matrix of each group. The Ksubspaces algorithm proceeds by alternating between these two steps until convergence to a local minimum.

In this section, we are interested in the more general problem of  $n > 1$  subspaces and  $m_j > 1$  centers per subspace. In principle, we could also solve this problem using Kmeans by interpreting Problem 1 as a central clustering problem with  $\sum m_j$  cluster centers. However, Kmeans does not fully employ the data's structural information and can cause undesirable clustering results, as shown in Figure 1.2. Thus, we propose a new algorithm which combines the objective functions (1.1) and (1.2) into a single objective. The algorithm is a natural generalization of both Kmeans and Ksubspaces to simultaneous central-subspace clustering.

For the sake of simplicity, let us first assume that the subspaces are of co-dimension one, i.e. hyperplanes, so that we can represent them with a single normal vector  $\mathbf{b}_j \in \mathbb{R}^D$ . We discuss the extension to subspaces of varying dimensions in Remark 1. Our method computes the cluster centers and the subspace normals by

solving the following optimization problem

$$\min \sum_{i=1}^P \sum_{j=1}^n \sum_{k=1}^{m_j} w_{ijk} ((\mathbf{b}_j^\top \mathbf{x}_i)^2 + \|\mathbf{x}_i - \mu_{jk}\|^2) \quad (1.3)$$

$$\text{subject to } \mathbf{b}_j^\top \mathbf{b}_j = 1, j = 1, \dots, n, \quad (1.4)$$

$$\mathbf{b}_j^\top \mu_{jk} = 0, j = 1, \dots, n, k = 1, \dots, m_j, \quad (1.5)$$

$$\sum_{j=1}^n \sum_{k=1}^{m_j} w_{ijk} = 1, i = 1, \dots, P, \quad (1.6)$$

where  $w_{ijk} \in \{0, 1\}$  denotes the membership of the  $i$ th point to the  $jk$ th cluster center. Equation (1.3) ensures that for each point  $\mathbf{x}_i$ , there is a subspace  $j$  and a cluster  $k$  such that both  $|\mathbf{b}_j^\top \mathbf{x}_i|$  and  $\|\mathbf{x}_i - \mu_{jk}\|$  are small. Equation (1.4) ensures that the normal vectors are of unit norm. Equation (1.5) ensures that each cluster center lies in its corresponding hyperplane and equation (1.6) ensures that each point is assigned to only one of the  $\sum m_j$  cluster centers.

Using the technique of Lagrange multipliers to minimize the cost function in (1.3) subject to the constraints (1.4)–(1.6) leads to the new objective function

$$\begin{aligned} \mathcal{L} = & \sum_{i=1}^P \sum_{j=1}^n \sum_{k=1}^{m_j} w_{ijk} ((\mathbf{b}_j^\top \mathbf{x}_i)^2 + \|\mathbf{x}_i - \mu_{jk}\|^2) + \\ & \sum_{j=1}^n \sum_{k=1}^{m_j} \lambda_{jk} (\mathbf{b}_j^\top \mu_{jk}) + \sum_{j=1}^n \delta_j (\mathbf{b}_j^\top \mathbf{b}_j - 1). \end{aligned} \quad (1.7)$$

Similarly to the Kmeans and Ksubspaces algorithms, we minimize  $\mathcal{L}$  using a coordinate descent minimization technique, as shown in Algorithm 1. The following subsections describe each step of the algorithm in detail.

**Initialization:** Since the data points lie in a collection of hyperplanes, we can apply GPCA to obtain an estimate of the normal vectors  $\{\mathbf{b}_j\}_{j=1}^n$  and segment the data into  $n$  groups. Let  $\mathbf{X}_j \in \mathbb{R}^{D \times P_j}$  be the set of points in the  $j$ th hyperplane. If we use the SVD of  $\mathbf{X}_j$  to compute a rank  $D - 1$  approximation of  $\mathbf{X}_j \approx U_j S_j V_j$ , where  $U_j \in \mathbb{R}^{D \times (D-1)}$ ,  $S_j \in \mathbb{R}^{(D-1) \times (D-1)}$  and  $V_j \in \mathbb{R}^{(D-1) \times P_j}$ , then the columns of  $\mathbf{X}'_j = S_j V_j \in \mathbb{R}^{(D-1) \times P_j}$  are a set of vectors in  $\mathbb{R}^{D-1}$  distributed around  $m_j$  cluster centers. We can apply Kmeans to segment the columns of  $\mathbf{X}'_j$  into  $m_j$  groups and obtain the projected cluster centers  $\{\mu'_{jk} \in \mathbb{R}^{D-1}\}_{k=1}^{m_j}$ . The original cluster centers are then given by  $\mu_{jk} = U_j \mu'_{jk} \in \mathbb{R}^D$ .

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**Algorithm 1** (Combined Central and Subspace Clustering)

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1. *Initialization:* Obtain an initial estimate of the normal vectors  $\{\mathbf{b}_j\}_{j=1}^n$  and cluster centers  $\{\mu_{jk}\}_{j=1 \dots n}^{k=1 \dots m_j}$  using GPCA followed by Kmeans in each subspace.
  2. *Computing the memberships:* Given the normal vectors  $\{\mathbf{b}_j\}_{j=1}^n$  and the cluster centers  $\{\mu_{jk}\}_{j=1 \dots n}^{k=1 \dots m_j}$ , compute the memberships  $\{w_{ijk}\}$ .
  3. *Computing the cluster centers:* Given the memberships  $\{w_{ijk}\}$  and the normal vectors  $\{\mathbf{b}_j\}_{j=1}^n$ , compute the cluster centers  $\{\mu_{jk}\}_{j=1 \dots n}^{k=1 \dots m_j}$ .
  4. *Computing the normal vectors:* Given the memberships  $\{w_{ijk}\}$  and the cluster centers  $\{\mu_{jk}\}_{j=1 \dots n}^{k=1 \dots m_j}$ , compute the normal vectors  $\{\mathbf{b}_j\}_{j=1}^n$ .
  5. *Iterate:* Repeat steps 2,3,4 until convergence of the memberships.
- 

**Computing the memberships:** Since the cost function is positive and linear in  $w_{ijk}$ , the minimum is attained at  $w_{ijk}=0$ . However, since  $\sum_{jk} w_{ijk}=1$ , the  $w_{ijk}$  multiplying the smallest  $((\mathbf{b}_j^\top \mathbf{x}_i)^2 + \|\mathbf{x}_i - \mu_{jk}\|^2)$  must be 1. Thus,

$$w_{ijk} = \begin{cases} 1 & \text{if } (j, k) = \arg \min ((\mathbf{b}_j^\top \mathbf{x}_i)^2 + \|\mathbf{x}_i - \mu_{jk}\|^2) \\ 0 & \text{otherwise} \end{cases}.$$

**Computing the cluster centers:** From the first order condition for a minimum we have

$$\frac{\partial \mathcal{L}}{\partial \mu_{jk}} = -2 \sum_{i=1}^P w_{ijk} (\mathbf{x}_i - \mu_{jk}) + \lambda_{jk} \mathbf{b}_j = 0. \quad (1.8)$$

Left-multiplying (1.8) by  $\mathbf{b}_j^\top$  and recalling that  $\mathbf{b}_j^\top \mu_{jk} = 0$  and  $\mathbf{b}_j^\top \mathbf{b}_j = 1$  yields

$$\lambda_{jk} = 2 \sum_{i=1}^P w_{ijk} (\mathbf{b}_j^\top \mathbf{x}_i). \quad (1.9)$$

Substituting (1.9) into (1.8) and dividing by two yields

$$\begin{aligned} - \sum_{i=1}^P w_{ijk} (\mathbf{x}_i - \mu_{jk}) + \sum_{i=1}^P w_{ijk} \mathbf{b}_j \mathbf{b}_j^\top \mathbf{x}_i &= 0 \\ \implies \boxed{\mu_{jk} = (\mathcal{I} - \mathbf{b}_j \mathbf{b}_j^\top) \frac{\sum_{i=1}^P w_{ijk} \mathbf{x}_i}{\sum_{i=1}^P w_{ijk}}} \end{aligned}$$

where  $\mathcal{I}$  is the identity matrix in  $\mathbb{R}^D$ . Note that the optimal  $\mu_{jk}$  has a simple geometric interpretation: it is the mean of the points associated with the  $jk$ th cluster, projected onto the  $j$ th hyperplane.

**Computing the normal vectors:** From the first order condition for a minimum we have

$$\frac{\partial \mathcal{L}}{\partial \mathbf{b}_j} = 2 \sum_{i=1}^P \sum_{k=1}^{m_j} w_{ijk} (\mathbf{b}_j^\top \mathbf{x}_i) \mathbf{x}_i + \sum_{k=1}^{m_j} \lambda_{jk} \mu_{jk} + 2\delta_j \mathbf{b}_j = 0. \quad (1.10)$$

After left-multiplying (1.10) by  $\mathbf{b}_j^\top$  to eliminate  $\lambda_{jk}$  and recalling that  $\mathbf{b}_j^\top \mu_{jk} = 0$ , we obtain

$$\delta_j = - \sum_{i=1}^P \sum_{k=1}^{m_j} w_{ijk} (\mathbf{b}_j^\top \mathbf{x}_i)^2. \quad (1.11)$$

After substituting (1.9) into equation (1.10) and recalling that  $\mathbf{b}_j^\top \mu_{jk} = 0$ , we obtain

$$(\sum_{i=1}^P \sum_{k=1}^{m_j} w_{ijk} (\mathbf{x}_i + \mu_{jk}) \mathbf{x}_i^\top + \delta_j \mathcal{I}) \mathbf{b}_j = 0. \quad (1.12)$$

Therefore, the optimal normal vector  $\mathbf{b}_j$  is the eigenvector of  $(\sum_{i=1}^P \sum_{k=1}^{m_j} w_{ijk} (\mathbf{x}_i + \mu_{jk}) \mathbf{x}_i^\top + \delta_j \mathcal{I})$  associated with its smallest eigenvalue, which can be computed via SVD.

**Remark 1** (Extension from hyperplanes to subspaces). *In the case of subspaces of co-dimension larger than one, each normal vector  $\mathbf{b}_j$  should be replaced by a matrix of normal vectors  $B_j \in \mathbb{R}^{D \times (D-d_j)}$ , where  $d_j$  is the dimension of the  $j$ th subspace. Since the normal bases and the means must satisfy  $B_j^\top \mu_{jk} = 0$  and  $B_j^\top B_j = \mathcal{I}$ , the objective function (1.3) should be changed to*

$$\begin{aligned} \mathcal{L} = & \sum_{i=1}^P \sum_{j=1}^n \sum_{k=1}^{m_j} w_{ijk} (\|B_j^\top \mathbf{x}_i\|^2 + \|\mathbf{x}_i - \mu_{jk}\|^2) + \\ & \sum_{j=1}^n \sum_{k=1}^{m_j} \lambda_{jk}^\top (B_j^\top \mu_{jk}) + \sum_{j=1}^n \text{trace}(\Delta_j (B_j^\top B_j - \mathcal{I})). \end{aligned}$$

where  $\lambda_{jk} \in \mathbb{R}^{(D-d_j)}$  and  $\Delta_j \in \mathbb{R}^{(D-d_j) \times (D-d_j)}$  are, respectively, vectors and matrices of Lagrange multipliers. Given the normal basis  $B_j$ , the optimal solution for the means is given by

$$\mu_{jk} = (\mathcal{I} - B_j B_j^\top) \frac{\sum_{i=1}^P w_{ijk} \mathbf{x}_i}{\sum_{i=1}^P w_{ijk}}.$$

One can show that the optimal solution for  $\Delta_j$  is a scaled identity matrix whose  $j$ th diagonal entry is  $\delta_j = - \sum_{i=1}^P \sum_{k=1}^{m_j} w_{ijk} \|B_j^\top \mathbf{x}_i\|^2$ . Given  $\delta_j$  and  $\mu_{jk}$ , one can still solve for  $B_j$  from the null space of  $(\sum_{i=1}^P \sum_{k=1}^{m_j} w_{ijk} (\mathbf{x}_i + \mu_{jk}) \mathbf{x}_i^\top + \delta_j \mathcal{I})$ , which can be proved to have dimension  $D - d_j$ .

**Remark 2** (Maximum Likelihood Solution). Notice that in the combined objective function (1.7) the term  $|\mathbf{b}_j^\top \mathbf{x}_i|$  is the distance to the  $j$ th hyperplane, while  $\|\mathbf{x}_i - \mu_{jk}\|$  is the distance to the  $jk$ th cluster center. Since the former is mostly related to the variance of the noise in the orthogonal direction to the hyperplane,  $\sigma_b^2$ , while the latter is mostly related to the within class variance,  $\sigma_\mu^2$ , the relative magnitudes of these two distances need to be taken into account. One way of doing so is to assume that the data is generated by a mixture of  $\sum m_j$  Gaussians with means  $\mu_{jk}$  and covariances  $\Sigma_{jk} = \sigma_b^2 \mathbf{b}_j \mathbf{b}_j^\top + \sigma_u^2 (\mathcal{I} - \mathbf{b}_j \mathbf{b}_j^\top)$ . This automatically allows the variances inside and orthogonal to the hyperplanes to be different. Application of the EM algorithm to this mixture model leads to the minimization of the following normalized objective function

$$\begin{aligned} \mathcal{L} = & \sum_{i=1}^P \sum_{j=1}^n \sum_{k=1}^{m_j} w_{ijk} \left( \frac{(\mathbf{b}_j^\top \mathbf{x}_i)^2}{2\sigma^2} + \frac{\|\mathbf{x}_i - \mu_{jk}\|^2}{2\sigma_\mu^2} + \log(\sigma_b) + \right. \\ & \left. (D-1) \log(\sigma_u) \right) + \sum_{j=1}^n \sum_{k=1}^{m_j} \lambda_{jk} (\mathbf{b}_j^\top \mu_{jk}) + \sum_{j=1}^n \delta_j (\mathbf{b}_j^\top \mathbf{b}_j - 1) \end{aligned}$$

where  $w_{ijk} \propto \exp(-\frac{(\mathbf{b}_j^\top \mathbf{x}_i)^2}{2\sigma^2} - \frac{\|\mathbf{x}_i - \mu_{jk}\|^2}{2\sigma_\mu^2})$  is now the probability that the  $i$ th point belongs to the  $jk$ th cluster center, and  $\sigma^{-2} = \sigma_b^{-2} - \sigma_\mu^{-2}$ . The optimal solution can be obtained using coordinate descent, similarly to Algorithm 1, as follows

$$\begin{aligned} \lambda_{jk} &= 2 \sum_{i=1}^P w_{ijk} \frac{\mathbf{b}_j^\top \mathbf{x}_i}{\sigma_\mu^2}, \quad \delta_j = - \sum_{i=1}^P \sum_{k=1}^{m_j} w_{ijk} \frac{(\mathbf{b}_j^\top \mathbf{x}_i)^2}{\sigma^2} \\ \mu_{jk} &= (\mathcal{I} - \mathbf{b}_j \mathbf{b}_j^\top) \frac{\sum_{i=1}^P w_{ijk} \mathbf{x}_i}{\sum_{i=1}^P w_{ijk}} \\ 0 &= \left( \sum_{i=1}^P \sum_{k=1}^{m_j} w_{ijk} \left( \frac{\mathbf{x}_i}{\sigma^2} + \frac{\mu_{jk}}{\sigma_\mu^2} \right) \mathbf{x}_i^\top + \delta_j \mathcal{I} \right) \mathbf{b}_j \\ \sigma_b^2 &= \frac{\sum_{i=1}^P \sum_{j=1}^n \sum_{k=1}^{m_j} w_{ijk} (\mathbf{b}_j^\top \mathbf{x}_i)^2}{\sum_{i=1}^P \sum_{j=1}^n \sum_{k=1}^{m_j} w_{ijk}} \\ \sigma_u^2 &= \frac{\sum_{ijk} w_{ijk} (\|\mathbf{x}_i - \mu_{jk}\|^2 - (\mathbf{b}_j^\top \mathbf{x}_i)^2)}{(D-1) \sum_{i=1}^P \sum_{j=1}^n \sum_{k=1}^{m_j} w_{ijk}}. \end{aligned}$$

## 1.3 Experiments: Clustering Performance Evaluation

### 1.3.1 Comparison: Central clustering and subspace clustering

Given a collection of data samples, the task is to group them into  $K$  clusters.

**Kmeans:** Kmeans algorithm first randomly selects K data samples as the seeds for each cluster. Then other data samples are assigned to the nearest-distanced seed. After this, the seeds of each clusters are replaced by the mean of all data samples that belong to the according cluster. The iterative procedure updates the data sample memberships and the cluster means alternatively until convergency.

**EM:** Expectation Maximization method can be considered as a soft-constrained Kmeans. The membership of a data sample to any cluster is not a binary decision as Kmeans, but rather a continuous value of probability. This probability measures the likelihood of the data sample with respective to the cluster model. EM algorithm depends on random initialization and iteratively converges to local maxima.

**Mixtures of Probabilistic PCA:** Mixtures of PPCA (MPPCA) [23] is a probabilistic generalization of principal component analysis by introducing latent random variables. There is an analytic solution for parameter estimation of PPCA given labeled data, but the data clustering process is still an iterative optimization by a EM fashion.

**K-subspace:** Given an initial estimate for the subspace bases, this algorithm alternates between clustering the data points using the distance residual to the different subspaces, and computing a basis for each subspace using standard PCA. See [8] for further details.

**Generalized PCA:** An algebraic solution for one-shot subspace clustering is recently proposed which is named Generalized Principal Component Analysis (GPCA) [25]. The union of subspaces is modeled with a homogeneous polynomial via the Veronese embedding. All data samples fit into the same polynomial function. The clustering problem then is solved by polynomial differentiation and division [25]. Each subspace is uniquely represented by the normal vector of the hyperplane.

Note that the clustering algorithms based on random initialization, eg. Kmeans, EM, MPPCA, Ksubspaces, converge to different local maximums with each start. Therefore they can be improved by multiple restarts and selecting the results with the best data-model fitness. On the other hand, the random initialization process is completely avoided by the combined constrained polynomial function and its algebraic solution in GPCA.

### 1.3.2 Performance Evaluation on simulated data

We randomly generate  $P = 600$  data points in  $\mathbb{R}^3$  lying in 2 intersecting planes  $\{S_j\}_{j=1}^2$  with 3 clusters in each plane  $\{\mu_{jk}\}_{j=1,2}^{k=1,2,3}$ . 100 points are drawn around each one of the six cluster centers according to a zero-mean Gaussian distribution with standard deviation  $\sigma_\mu = 1.5$  within each plane. The angle between the two planes is randomly chosen from  $20^\circ \sim 90^\circ$ , and the distance among the three cluster centers is randomly selected in the range  $2.5\sigma_\mu \sim 5\sigma_\mu$ . Zero-mean Gaussian noise with standard deviation  $\sigma_b$  is added in the direction orthogonal to each plane. Using simulated data, we compare 5 different clustering methods:

- Kmeans clustering in  $\mathbb{R}^3$  using 6 cluster centers, then merging them into 2 planes<sup>2</sup> (**KM**),
- MPPCA<sup>3</sup> clustering in  $\mathbb{R}^3$  using 6 cluster centers, then merging them into 2 planes<sup>1</sup> (**MP**),
- Ksubspaces clustering in  $\mathbb{R}^3$  using 2 planes, then Kmeans using 3 clusters within each plane (**KK**),
- GPCA clustering in  $\mathbb{R}^3$  using 2 planes, then Kmeans using 3 clusters within each plane (**GK**),
- GPCA-Kmeans clustering for initialization followed by combined central and subspace clustering (**JC**) as described in Section 1.2 (Algorithm 1).

Figure 1.3 shows a comparison of the performance of these five methods in terms of clustering error ratios and the error in the estimation of the subspace normals in degrees. The results are the mean of the errors over 100 trials. It can be seen in Figure 1.3 that the errors in clustering and normal vectors of all five algorithms increase as a function of noise. **MP** performs better than **KM** and **KK** for large levels of noise, because of its probabilistic formulation. The two stage algorithms, **KK**, **GK** and **JC**, in general perform better than **KM** and **MP** in terms of clustering error. The random initialization based methods, **KM**, **MP** and **KK**, have non-zero clustering error even with noise-free data. Within the two stage algorithms, **KK**

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<sup>2</sup>In order to estimate the plane normals, we group the 6 clusters returned by **KM** or **MP** into 2 planes. The idea is that 3 clusters which lie in the same plane have the dimensionality of 2 instead of 3. A brute-force search with  $\binom{6}{3}/2$  selections is employed to find the 2 best fitting planes, by considering the minimal strength of the data distributed in the third dimension via Singular Value Decomposition [5].

<sup>3</sup>Software available at [www.ncrg.aston.ac.uk/netlab/](http://www.ncrg.aston.ac.uk/netlab/)

begins to experience subspace clustering failures more frequently with more severe noises, due to its random initialization, while GPCA in **GK** and **JC** employ an algebraic solution of one-shot subspace clustering, thus avoiding the initialization problem. The subspace clustering errors of **KK** can cause the estimate of the normals to be very inaccurate, which explains why **KK** has worse errors in the normal vectors than **KM** and **MP**. In summary, **GK** and **JC** have smaller average errors in clustering and normal vectors than **KM**, **MP** and **KK**. The combined optimization procedure of **JC** converges within  $2 \sim 5$  iterations according to our experiments, which further advocates **JC**'s clustering performance.

## 1.4 Experiments: Illumination-invariant face clustering

The Yale face database B (see <http://cvc.yale.edu/projects/yalefacesB/yalefacesB.html>) contains a collection of face images  $I_j \in \mathbb{R}^K$  of 10 subjects taken under 576 viewing conditions (9 poses  $\times$  64 illumination conditions). Here we only consider the illumination variation for face clustering in the case of frontal face images. Thus our task is to sort the images taken for the same person by using our combined central-subspace clustering algorithm. As shown in [8], the set of all images of a (Lambertian) human face with fixed pose taken under all lighting conditions forms a cone in the image space which can be well approximated by a low dimensional subspace. Thus images of different subjects live in different subspaces. Since the number of pixels  $K$  of each image is in general much larger than the dimension of the underlying subspace, PCA [5] is first employed for dimensionality reduction. Successful GPCA clustering results have been reported by [25] for a subset of 3x64 images of subjects 5, 8 and 10. The images in [25] are cropped to 30x40 pixels and 3 PCA components are used as image features in homogeneous coordinates.

In this subsection, we further explore the performance of combined central-subspace face clustering under more complex imaging conditions. We keep 3 PCA components for 4x64 (240x320 pixels) images of subjects 5, 6, 7, and 8, which gives more background details (as shown in Figure 1.4). Figures 1.5 (a,b) show the imperfect clustering result of GPCA due to the intersection of the subspace of subject 5 with the subspaces of subjects 6 and 7. GPCA assigns all the images on the intersection to subject 5. Mixtures of PPCA is implemented in Netlab as a probabilistic variation of subspace clustering with one spatial center per

subspace. It can be initialized with Kmeans (originally in Netlab) or GPCA, both of which result in imperfect clustering. We show one example of the subspaces of subjects 6 and 7 mixed (Kmeans initialization) in Figure 1.5 (c,d). Our combined subspace-central optimization process successfully corrects the wrong labels for some images of subjects 6 and 7, as demonstrated in Figure 1.5 (e,f). In the optimization, the local clusters in the subspaces of subjects 6 and 7 contribute with smaller central distances to their misclassified images, which re-classifies them to the correct subspaces using our combined subspace-central clustering algorithm. In this experiment, 4 subspaces with 2 clusters per subspace are used. Compared with the results in [25], we obtain perfect illumination-invariant face clustering for a more complex data distribution.

## 1.5 Experiments: Video shot segmentation

Unlike face images under different illumination conditions, video data provides continuous visual signals. Video structure parsing and analysis applications need to segment the whole video sequence into several video shots. Each video shot may contain hundreds of image frames which are either captured with a similar background or have a similar semantical meaning.

Figure 1.6 shows 2 sample videos, *mountain.avi* and *drama.avi*, containing 4 shots each. Archives are publicly available from <http://www.open-video.org>. For the mountain sequence, 4 shots are captured. The shots display different backgrounds and show either multiple dynamic objects and/or severe camera motions. In this video, the frames between each pair of successive shots are gradually blended from one to another. Because of this, the correct video shot segmentation is considered to split every two successive shots at their blending frames. In order to explore how the video frames are distributed in feature space, we plot the first 3 PCA components for each frame in Figure 1.7 (b, d, f). Note that a manifold structure can be observed in Figure 1.7 (f), where we manually label each portion of the data as shots 1 through 4 (starting from red dots to green, black and ending in blue) according to the result of our clustering method. The video shot segmentation results of the mountain sequence by Kmeans, GPCA and GPCA-Kmeans followed by combined optimization are shown in Figure 1.7 (a,b), (c,d) and (e,f), respectively. Because Kmeans is based on the central distances among data, it segments the data into spatially close blobs. There is no guarantee

that these spatial blobs will correspond to correct video shots. Comparing Figure 1.7 (b) with the correct segmentation in (f), the Kmeans algorithm splits shot 2 into clusters 2 and 3, while it groups shots 1 and 4 into cluster 1. By considering the data's manifold nature, GPCA provides a more effective approximation with multiple planes to the manifold in  $\mathbb{R}^3$  than the spatial blobs given by central clustering. The essential problem for GPCA is that it only deploys the co-planar condition in  $\mathbb{R}^3$ , without any constraint relying on their spatial locations. In the structural approximation of the data's manifold, there are many intersecting data points among 4 planes. These data points represent video frames with the clustering ambiguity solely based on the subspace constraint. Fortunately this limitation can be well tackled by GPCA-Kmeans with combined optimization. Combining central and subspace distances provides correct video shot clustering results for the mountain sequence, as demonstrated in Figure 1.7 (e,f).

The second video sequence shows a drama scenario which is captured with the same background. The video shots should be segmented by the semantic meaning of the performance of the actor and actress. In Figure 1.6 **Right**, we show 2 sample images for each shot. This drama video sequence contains very complex actor and actress' motions in front of a common background, which results in a more complex manifold data structure<sup>4</sup> than that of the mountain video. For better visualization, the normal vectors of data samples recovered by GPCA or the combined central-subspace optimization, are drawn originating from each data point in  $\mathbb{R}^3$  with different colors for each cluster. For this video, the combined optimization process shows a smoother clustering result in Figure 1.8 (c,d), compared with (a,b). In summary, GPCA can be considered as an effective way to group data in a manifold into multiple subspaces or planes in  $\mathbb{R}^3$  which normally better represent video shots than central clustering. GPCA-Kmeans with combined optimization can then associate the data at the intersection of planes into the correct clusters by optimizing combined distances. Subspace clustering seems to be a better method to group the data on a manifold by somehow preserving their geometric structure. Central clustering, such as Kmeans<sup>5</sup>, provides a piecewise constant approximation; while

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<sup>4</sup>Because there are image frames of transiting subject motions from one shot to another, the correct video shot segmentation is considered to split successive shots at their transiting frames.

<sup>5</sup>Due to space limitation, we do not provide the clustering result using Kmeans for this sequence which is similar with Figure 1.7 (a,b).

subspace clustering shows a piecewise linear approximation. On the other hand, subspace clustering can meet severe clustering ambiguity problems when the shape of the manifold is complex, as shown in Figure 1.8 (b,d). In this case, there are many intersections of subspaces so that subspace clustering results can be very sparse, without considering the spatial coherence. Combined optimization of central and subspace distances demonstrates superior clustering performance with real video sequences.

## 1.6 Discussion on Model Selection

Throughout the paper we have assumed that the number of subspaces  $n$ , their dimensions  $d_i$  and the number of clusters within each subspace  $m_j$  are known. In practice, these quantities may not be known beforehand.

When the number of subspaces is  $n = 1$ , the estimation of the dimension of the subspace  $d$  is essentially equivalent to the estimation of the number of principal components of the data set. This problem can be tackled by combining PCA with existing model selection techniques, such as minimum description length, Akaike information criterion, or Bayesian information criterion [5]. Given  $d$ , the number of clusters  $m$  can be determined by combining the Kmeans cost functional with the aforementioned model selection criteria.

When the number of subspaces is  $n > 1$ , one possible solution is to employ model selection algorithms for subspace and central clustering separately in a sequential manner, to determine  $n$  first, then  $d_i$  and then  $m_j$ . As shown in [25], GPCA provides a way of determining  $n$  from a rank constraint on a polynomial embedding of the data. Given  $n$ , one may cluster the data using GPCA, and then determine the dimension of each subspace as the number of principal components of the data points that belong to each subspace. Given  $n$  and  $d_i$ , one can use the model selection procedure mentioned earlier to determine the number of clusters  $m_j$  in Kmeans.

However, this three-stage solution is clearly not optimal. Ideally one would like to have a model selection criteria that integrates both types of clustering into one joint or combined process. This is obviously more difficult than combining the clustering algorithms, and is under current investigation.

### 1.6.1 Practical Solution

We address the model selection issue of our algorithm for real data in two cases. Firstly, if strictly following the definition of Problem 1, both the number of subspace  $n$  and the number of clusters within each subspace  $m$  need to be known beforehand our clustering algorithm. We can employ the model selection algorithms for subspace and central clustering separately in a sequential manner, to determine  $n$  first and then  $m$ . How to integrate the model selection criteria of two types of clustering into one joint or combined process is more difficult than the combination of the clustering algorithms itself, and is under investigation. The examples are toy problem 2 and multiple moving object clustering. The case is more like using additional subspace constraints to balance the central clustering results, which results in a combined distance metric. Secondly, for examples of toy problem 1, face clustering and video segmentation, they are essentially subspace clustering problems and only the number of subspace  $n$  turns out to be critical. This case is using (spatial) central distance constraints to solve the inherent ambiguity of subspace clustering at the intersection of any two subspaces. Empirically,  $m$  can vary within a moderate range ( $2 \sim 5$ , in our experiments) with similar clustering results.<sup>6</sup>

For the toy examples in Figure 1.1, face clustering 1.4 and temporal video segmentation 1.5, the number of clusters inside each subspace is one, thus only the number of subspace  $n$  turns out to be critical. Empirically,  $m$  can vary within a moderate range ( $2 \sim 5$ , in our experiments) with similar clustering results.

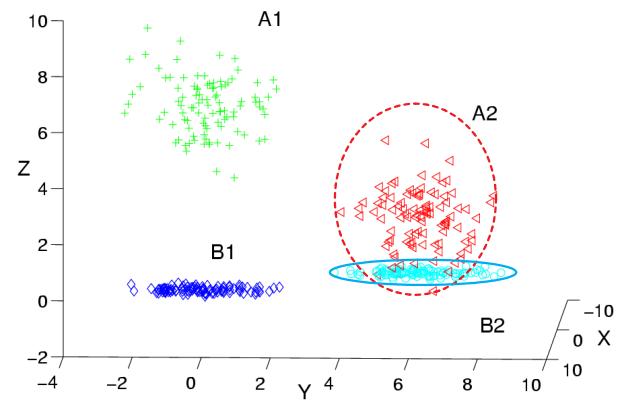
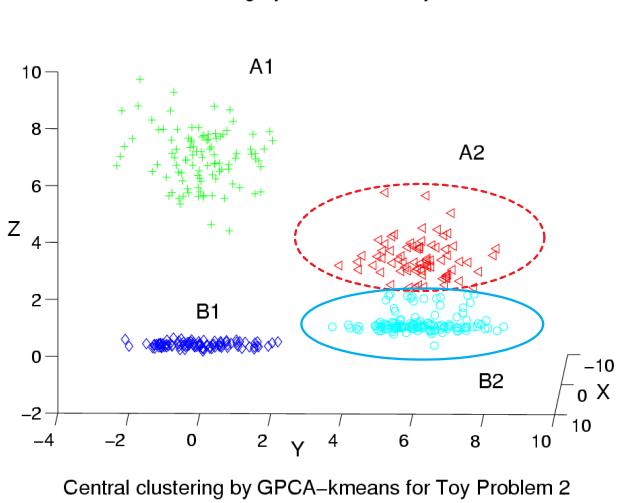
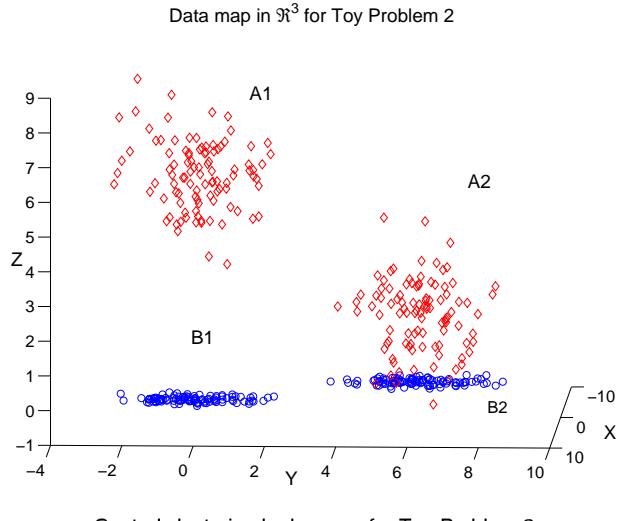
## 1.7 Conclusions and Future Work

We have proposed an intuitive and easy to implement algorithm for clustering data lying in a union of subspaces with multiple clusters within each subspace. By minimizing a cost function that incorporates both

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<sup>6</sup>If  $m$  is too large, the spatial constraints will be too local, which loses the stability of global data structure. For instance, the incorrectly subspace-clustered data samples in Figure 1.1 **center** can form a separate central cluster in the y-z plane, and the central constraints from the x-y plane become invalid. If  $m$  is too small, the spatial constraints will be too global so that the clustering algorithm can not be adaptive to local structures. Especially for loosely spatially distributed data inside each subspace, data samples which are too far should contribute less to the central based constraints on the samples near subspace intersections.

central and subspace distances, our algorithm can handle situations in which Kmeans and Ksubspaces/GPCA fail, e.g., when data are close to the intersection of two subspaces, or when cluster centers in different subspaces are spatially close. Future work includes using model selection to automatically determine the number of subspaces and cluster centers. Also, we believe it should be possible to extend the proposed combined central and subspace clustering formulation to recognize multiple complex curved manifolds. An example application is to find which movie a given images appear in. Each manifold will be composed of multiple subspaces where each subspace is spatially constrained by central distances among data samples. Once the movie models are learned (similarly to shot detection), the likelihood evaluation for a new data sample is based on computing its combined central and subspace distances to the given models.



**Figure 1.2:** **Top:** A set of points in  $\mathbb{R}^3$  distributed around 4 clusters labeled as  $A_1, A_2, B_1, B_2$ . Clusters  $B_1$  and  $B_2$  lie in the x-y plane and clusters  $A_1$  and  $A_2$  lie in the y-z plane. Note that cluster  $B_2$  (in blue) is spatially close to cluster  $A_2$  (in red). **Middle:** Central clustering by Kmeans assigns some points in  $A_2$  to  $B_2$ . **Bottom:** Subspace clustering using GPCA followed by central clustering inside each subspace using Kmeans gives the correct clustering into four groups.

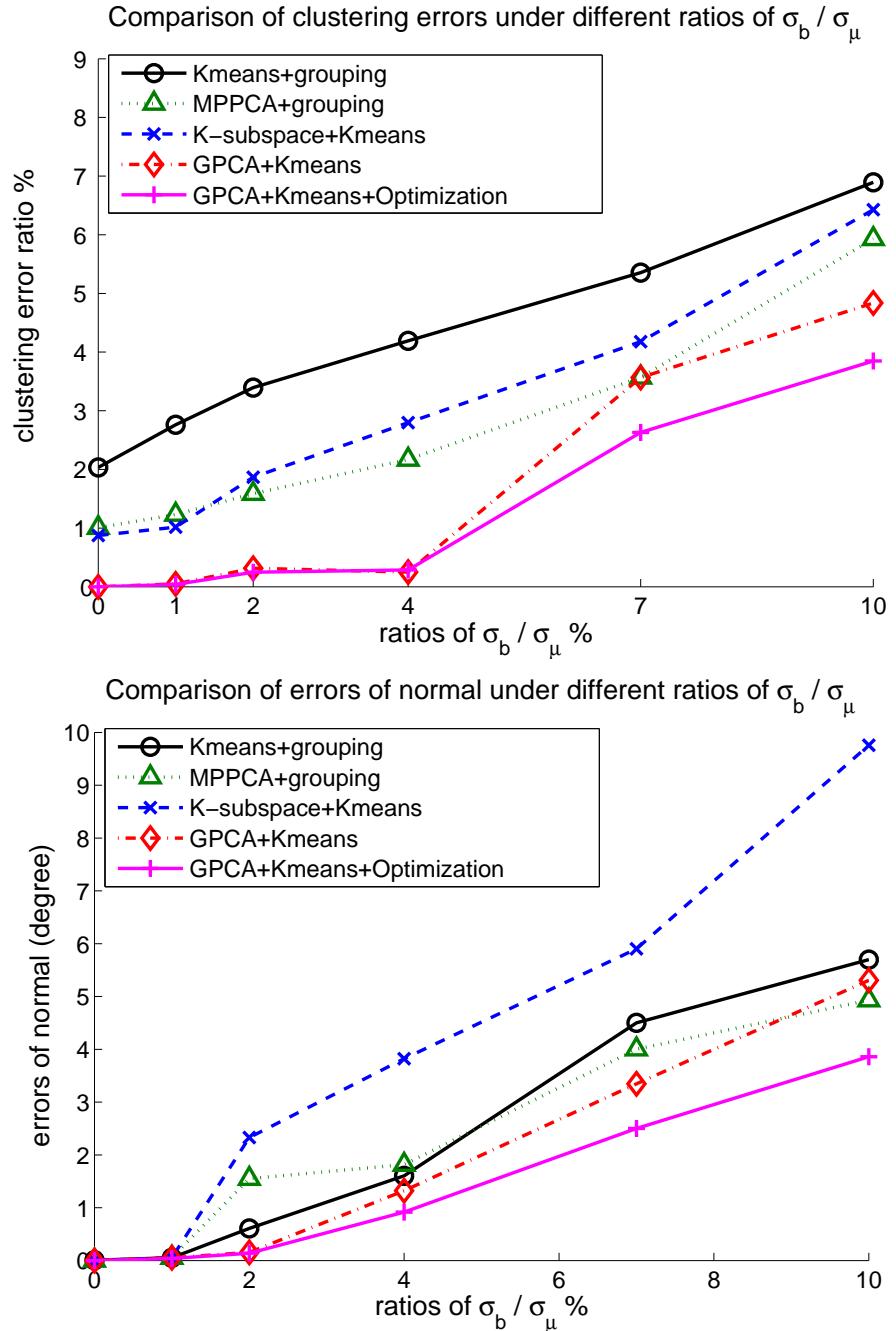


Figure 1.3: **Top:** Clustering error as a function of noise in the data. **Bottom:** Error in the estimation of the normal vectors (degrees) as a function of the level of noise in the data.

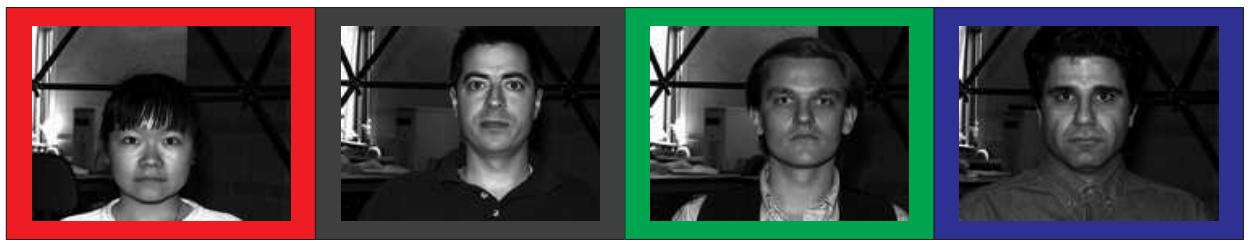


Figure 1.4: Sample images of subjects 5, 6, 7 and 8 shown in different colors.

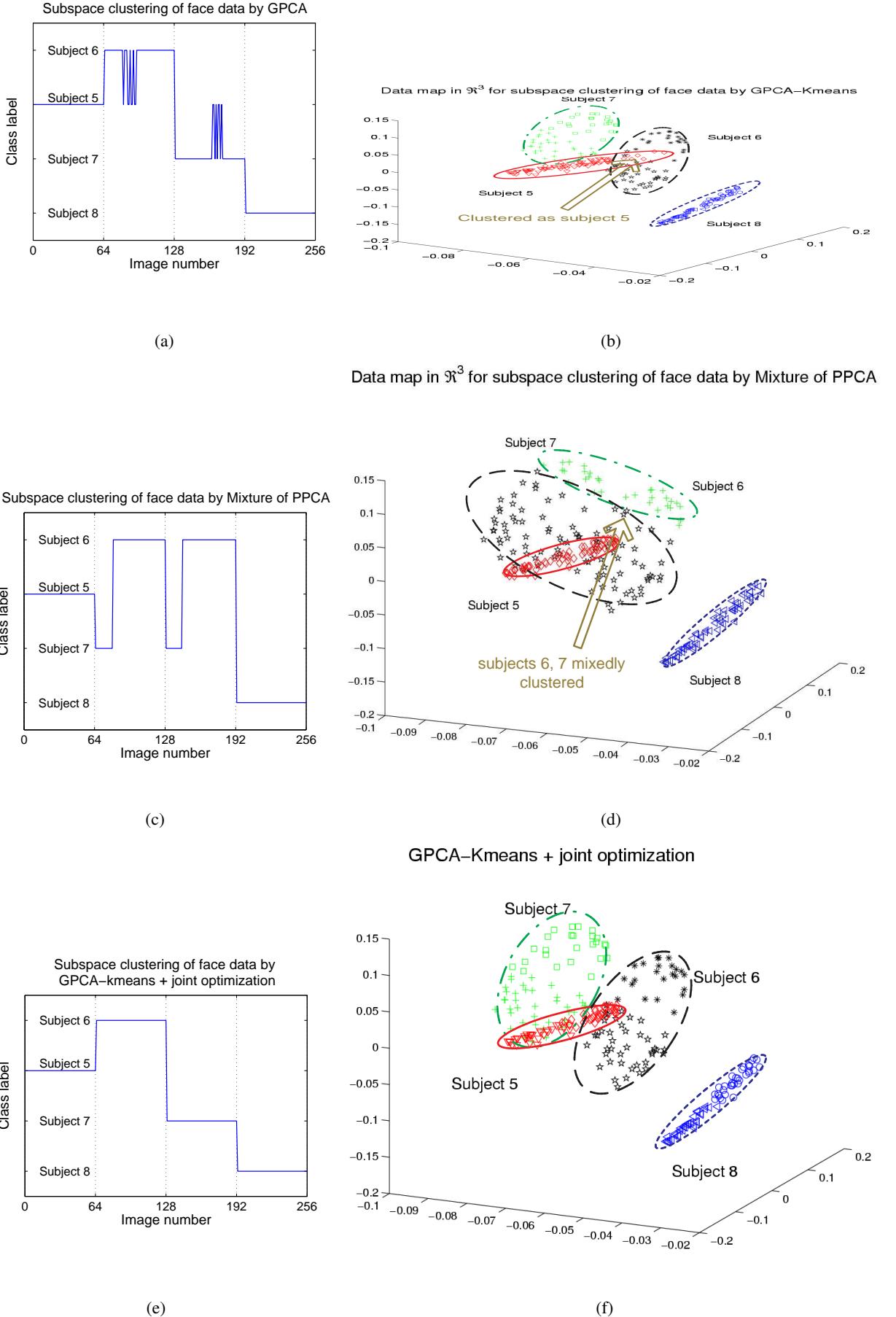


Figure 1.5: Illumination-invariant face clustering by GPCA (a-b), Mixtures of PPCA (c-d), and our method (e-f).

Plots on the right show 3 principal components with proper labels and color-shapes. The colors match the colors of subjects 5, 6, 7 and 8 in Figure 1.4.

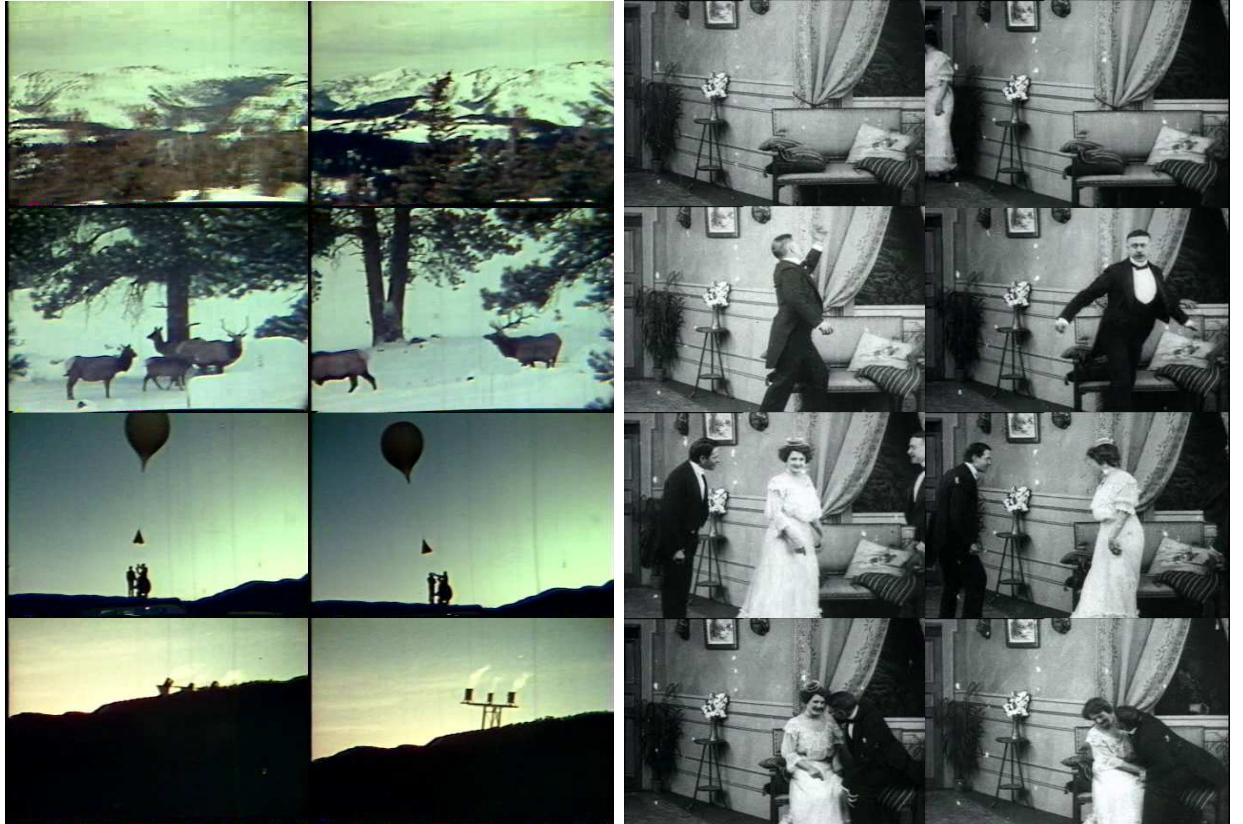


Figure 1.6: Sample images used for video shot segmentation. **Left:** Sample images from the mountain sequence. There are 4 shots in the video. Each row shows two images from each shot. All 4 shots are dynamic scenes, including large camera panning in shot 1, multiple animals moving in shot 2, a balloon rolling left and right in shot 3 and a rocket firing with the camera moving in shot 4. **Right:** Sample images from the drama sequence. There are 4 shots in the video. Each row shows 2 images from each shot. Shot 1 mainly shows the background only with no or little appearance of the actor or actress; shot 2 shows the actor's motion; shot 3 shows a scene of the actor and actress talking while standing; shot 4 shows the actor and actress kissing each other and sitting.

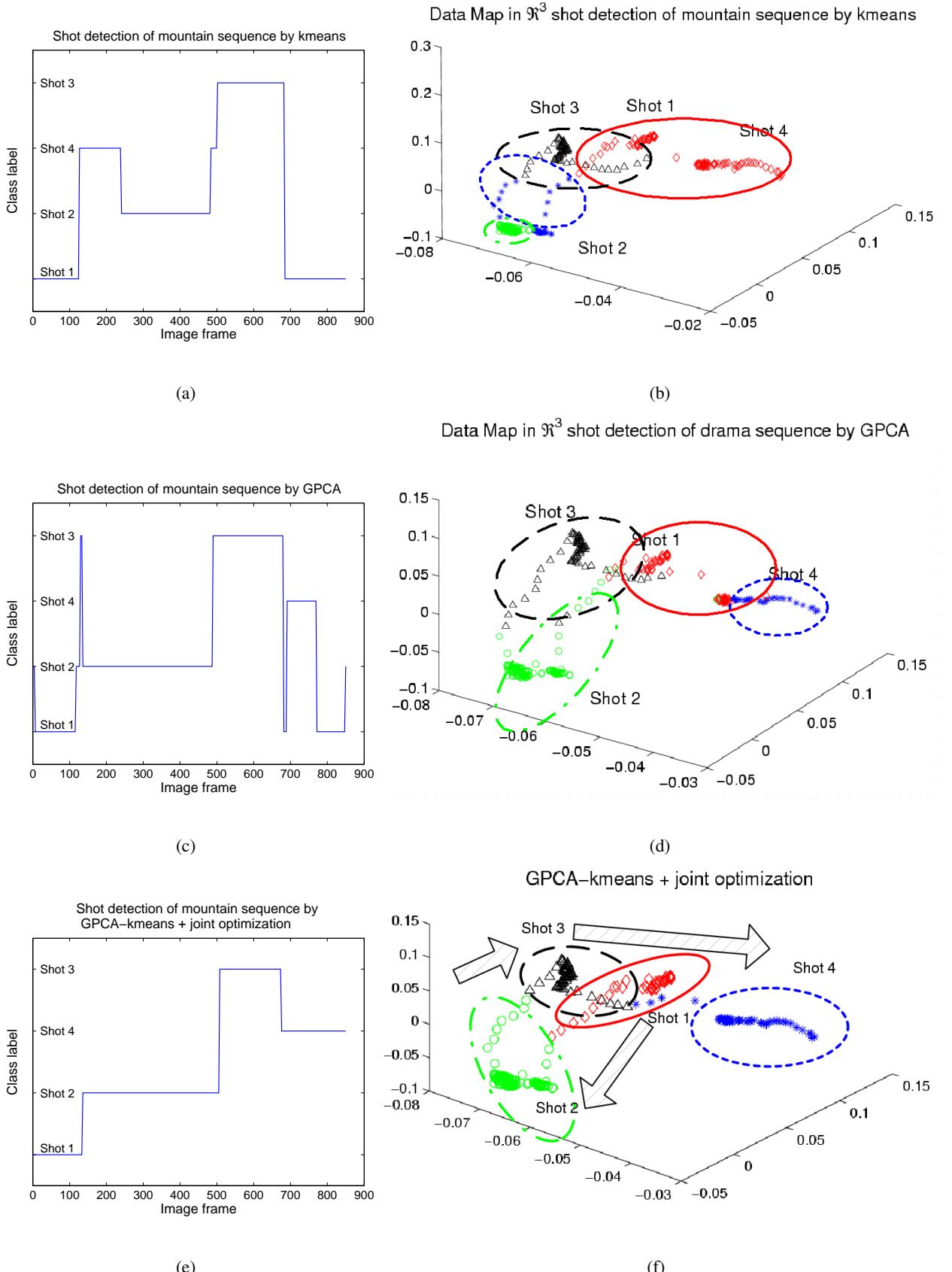


Figure 1.7: Video shot segmentation of mountain sequence by Kmeans (a-b), GPCA (c-d) and our algorithm (e-f). Plots on the right show 3 principal components of the data grouped in 4 clusters shown by ellipses with proper color-shapes. In (f), three arrows show the topology of the video manifold.

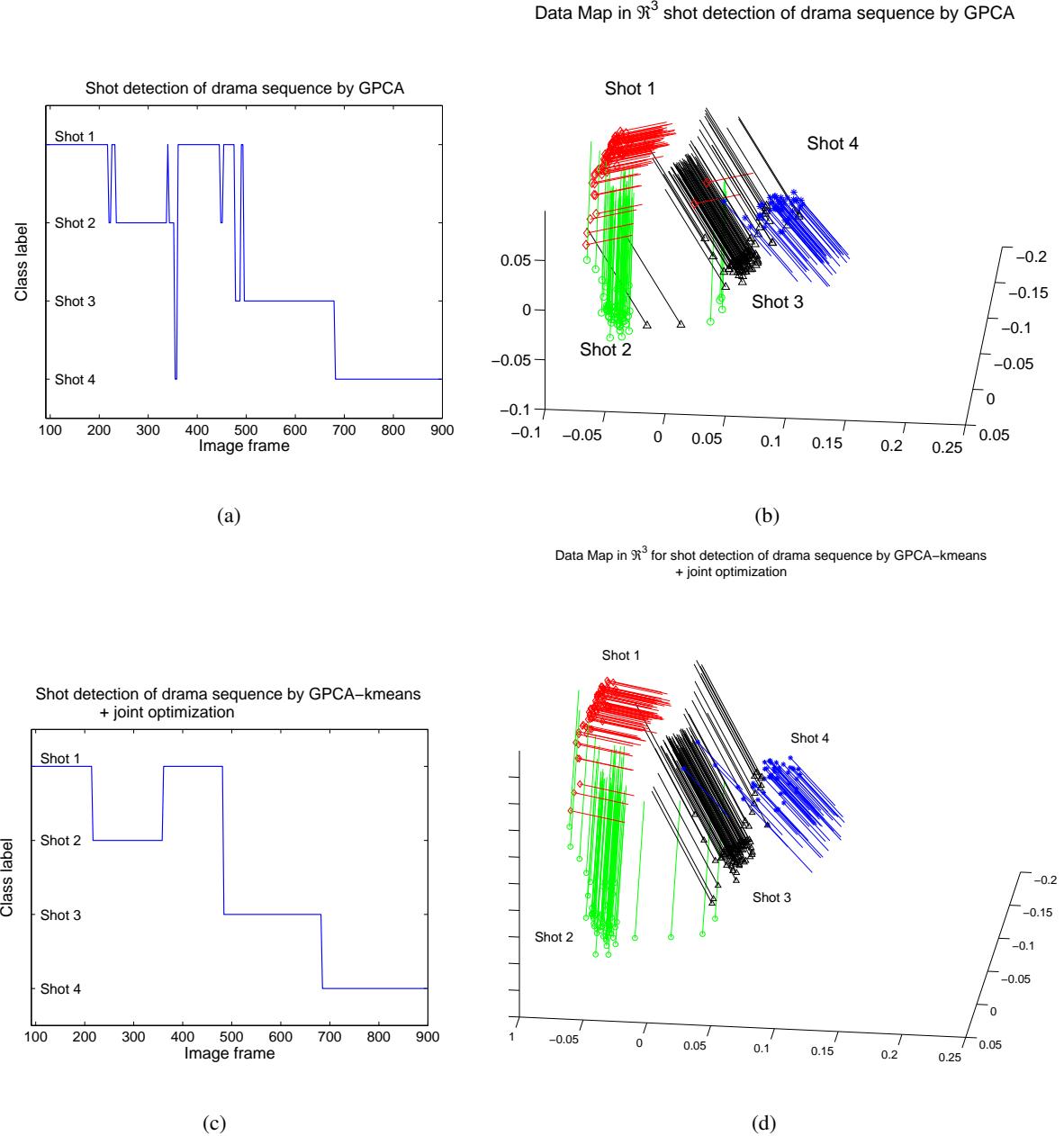


Figure 1.8: Video shot segmentation of drama sequence by GPCA (a-b), and our algorithm (c-d). Plots on the right show 3 principal components of the data with the normal to the plane at each point. Different normal directions illustrate different shots.

## Chapter 2

# Recognizing Temporal Units of Video using

## Textons:

### A Three-tiered Bottom-up Approach of

### Classifying Articulated Object Actions

*Visual action recognition is an important problem in computer vision. In this paper, we propose a new method to probabilistically model and recognize actions of articulated objects, such as hand or body gestures, in image sequences. Our method consists of three levels of representation. At the low level, we first extract a feature vector invariant to scale and in-plane rotation by using the Fourier transform of a circular spatial histogram. Then, spectral partitioning [135] is utilized to obtain an initial clustering; this clustering is then refined using a temporal smoothness constraint. Gaussian mixture model (GMM) based clustering and density estimation in the subspace of linear discriminant analysis (LDA) are then applied to thousands of image feature vectors to obtain an intermediate level representation. Finally, at the high level we build a temporal multi-resolution histogram model for each action by aggregating the clustering weights of sampled images belonging to that action. We discuss how this high level representation can be extended*



Figure 2.1: A gesture of finger spelling from 'I' to 'K', starting from frame 475# and ending at frame 499#.

*to achieve temporal scaling invariance and to include Bi-gram or Multi-gram transition information. Both image clustering and action recognition/segmentation results are given to show the validity of our three tiered representation.*

## 2.1 Introduction

Articulated object action modeling, tracking and recognition has been an important research issue in computer vision community for decades. Past approaches [35, 45, 36, 38, 56, 34] have used many different kinds of direct image observations, including color, edges, contour or moments [46], to fit a hand or body's shape model and motion parameters.

In this paper, we propose to learn a small set of object appearance descriptors, and then to build an aggregated temporal representation of clustered object descriptors over time. There are several obvious reasons to base gesture or motion recognition on a time sequence of observations. First, most hand or body postures are ambiguous. For example, in American Sign Language, 'D' and 'G', 'H' and 'U' have indistinguishable appearance from some viewpoints. Furthermore, these gestures are difficult to track from frame to frame due to motion blur, lack of features, and complex self-occlusions. An example of a less-than-one-second gesture of finger spelling from 'I' to 'K' is shown in Figure 2.1. By modeling hand/body gesture as a sequential learning problem, appropriate discriminative information can be retrieved and more action categories can be handled.

In related work, Darrell and Pentland [39] describe dynamic time warping (DTW) to align and recognize a space-time gesture against a stored library. To build the library, key views are selected from incoming an video by choosing views that have low correlation with all current views. This approach is empirical and

does not guarantee any sort of global consistency of the chosen views. As a result, recognition may be unstable. In comparision, our method describes image appearances uniformly and clusters them globally from a training set containing different gestures.

For static hand posture recognition, Tomasi et al. [57] apply vector quantization methods to cluster images of different postures and different viewpoints. This is a feature-based approach, with thousands of features extracted for each image. However, clustering in a high dimensional space is very difficult and can be unstable. We argue that fewer, more global features are adequate for the purposes of gesture recognition. Furthermore, the circular histogram representation has adjustable spatial resolution to accomodate differing appearance complexities, and it is translation, rotation, and scale invariant.

In other work, [60, 41] recognize human actions at a distance by computing motion information between images and relying on temporal correlation on motion vectors across sequences. Our work also makes use of motion information, but does not rely exclusively on it. Rather, we combine appearance and motion cues to increase sensitivity beyond what either can provide alone. Since our method is based on the temporal aggregation of image clusters as a histogram to recognize an action, it can also be considered to be a temporal texton-like method [121, 49]. One advantage of the aggregated histogram model in a time-series is that it is straightforward to accommodate temporal scaling by using a sliding window. In addition, higher order models corresponding to bigrams or trigrams of simpler “gestemes” can also be naturally employed to extend the descriptive power of the method.

In summary, there are four principal contributions in this paper. First, we propose a new scale/rotation-invariant hand image descriptor which is stable, compact and representative. Second, we introduce a methods for sequential smoothing of clustering results. Third, we show LDA/GMM with spectral partitioning initialization is an effective way to learn well-formed probability densities for clusters. Finally, we recognize image sequences as actions efficiently based on a flexible histogram model. We also discuss improvement to the method by incorporating motion information.

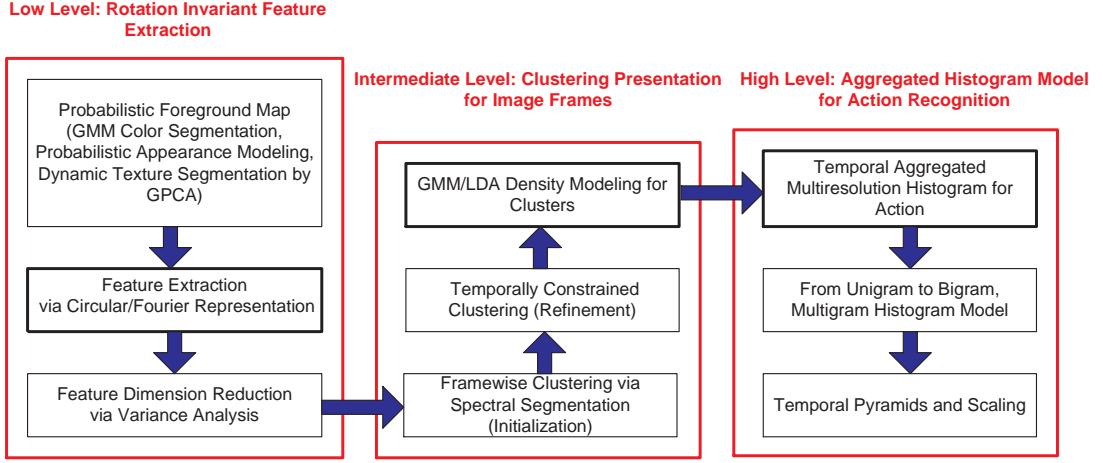


Figure 2.2: Diagram of a three tier approach for dynamic articulated object action modeling.

## 2.2 A Three Tiered Approach

We propose a three tiered approach for dynamic action modeling comprising low level feature extraction, intermediate level feature vector clustering and high level histogram recognition as shown in Figure 2.2.

### 2.2.1 Low Level: Rotation Invariant Feature Extraction

In the low level image processing, our goals are to locate the region of interest in an image and to extract a scale and in-plane rotation invariant feature vector as its descriptor. In order to accomplish this, a reliable and stable foreground model of the target in question is expected. Depending on the circumstances, a Gaussian mixture model (GMM) for segmentation [48], Maximum-Entropy color model ??, probabilistic appearance modeling [37], or dynamic object segmentation by Generalized Principal Component Analysis (GPCA) [58] are possible solutions. In this paper, we apply a GMM for hand skin color segmentation.

We fit a GMM by first performing a simple background subtraction to obtain a noisy foreground containing a hand object (shown in Figure 2.3 (a)). From this, more than 1 million RGB pixels are used to train skin and non-skin color density models with 10 Gaussian kernels for each class. Having done this, for new images a probability density ratio  $P_{skin}/P_{nonskin}$  of these two classes is computed. If  $P_{skin}/P_{nonskin}$  is larger than 1, the pixel is considered as skin (foreground) and is otherwise background. A morphological operator is then used to clean up this initial segmentation and create a binary mask for the hand object.

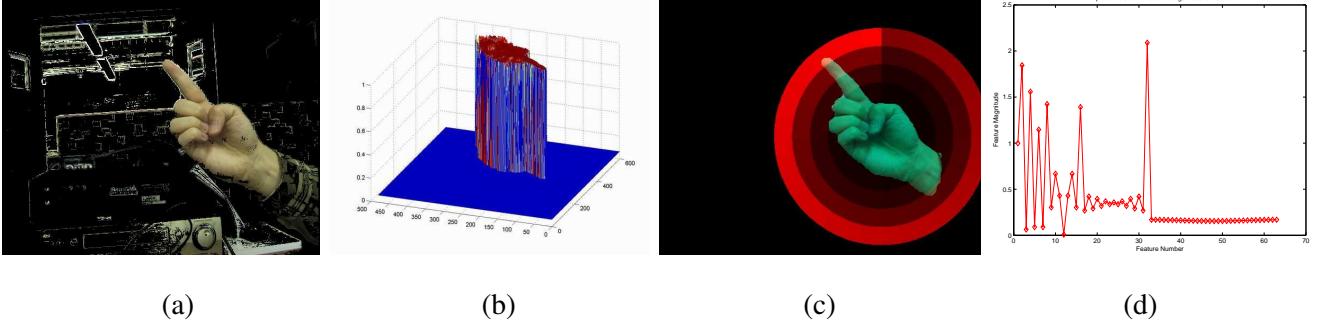


Figure 2.3: (a) Image after background subtraction (b) GMM based color segmentation (c) Circular histogram for feature extraction. (d) In-plane rotation invariant feature vector with 63 dimensions

We then compute the centroid and second central moments of this 2D mask. A circle is defined about the target by setting its center as the centroid and its radius as 2.8 times largest eigenvalues of the second central moment matrix (covering over 99% skin pixels in Figure 2.3 (c)). This circle is then divided to have 6 concentric annuli which contain 1, 2, 4, 8, 16, 32 bins from inner to outer, respectively. Since the position and size of this circular histogram is determined by the color segmentation, it is translation and scale invariant.

We then normalize the density value  $P_{skin} + P_{nonskin} = 1$  for every pixel within the foreground mask (Figure 2.3) over the hand region. For each bin of the circular histogram, we calculate the mean of  $P_{skin}$  ( $-\log(P_{skin})$ , or  $-\log(P_{skin}/P_{nonskin})$  are also possible choices) of pixels in that bin as its value. The values of all bins along each circle form a vector, and 1D Fourier transform is applied to this vector. The power spectra of all annuli are ordered into a linear list producing a feature vector  $\vec{f}(t)$  of 63 dimensions representing the appearance of a hand image.<sup>1</sup> Note that the use of the Fourier power spectrum of the annuli makes the representation rotation invariant.

### 2.2.2 Intermediate Level: Clustering Presentation (Textons) for Image Frames

After the low level processing, we obtain a scale and rotation invariant feature vector as an appearance representation for each image frame. The temporal evolution of feature vectors represent actions. However, not all the images are actually unique in appearance. At the intermediate level, we cluster images from a set of feature vectors. This frame-wise clustering is critical for dimension reduction and the stability of high

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<sup>1</sup>An optional dimension reduction of feature vectors can be achieved by eliminating dimensions which have low variance. It means that feature values of those dimensions do not change much in the data, therefore are non-informative.

level recognition.

### Initializing Clusters by Spectral Segmentation

There are two critical problems with clustering algorithms: determining the true number of clusters and initializing each cluster. Here we use a spectral clustering method [135, 147, 59, 51] to solve both problems. We first build the affinity matrix of pairwise distances between feature vectors<sup>2</sup>. We then perform a singular value decomposition on the affinity matrix with proper normalization [135]. The number of clusters is determined by choosing the  $n$  dominant eigenvalues. The corresponding eigenvectors are taken as an orthogonal subspace for all the data.

To get  $n$  cluster centers, we take the approach of [135] and choose vectors that minimize the absolute value of cosine between any two cluster centers:

$$ID(k) = \begin{cases} \text{rand}(0, N) & \text{if } k = 1; \\ \arg \min_{t=1..N} \sum_{c=1}^{k-1} |\cos(\vec{f}^n(ID(c)), \vec{f}^n(t))| & \text{if } n \geq k > 1; \end{cases} \quad (2.1)$$

where  $\vec{f}^n(t)$  is the feature vector of image frame  $t$  after numerical normalization in [135] and  $ID(k)$  is the image frame number chosen for the center of cluster  $k$ .  $N$  is the number of images used for spectral clustering. For better clustering results, multiple restarts are used for initialization.

Unlike [51], we find this simple clustering procedure is sufficient to obtain a good set of clusters from only a few restarts. After initialization, the Kmeans [5] is used to smooth the centers. Let  $C_1(t)$  denote the class label for image  $t$ , and  $\vec{g}(c) = \vec{f}(ID(c)); c = 1 \dots n$  denote cluster centers.

### Refinement: Temporally Constrained Clustering

Spectral clustering methods are designed for an unordered “bag” of feature vectors, but, in our case, the temporal ordering of image is an important source of information. In particular, the stability of appearance is

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<sup>2</sup>The exponent of either Euclidean distance or Cosine distance between two feature vectors can be used in this case.

easily computed by computing the motion energy<sup>3</sup> between two frames. Let  $E(t)$  denote the motion energy between frames  $t$  and  $t-1$ . Define  $T_{k,j} = \{t | C_1(t) = k, C_1(t-1) = j\}$  and  $\bar{E}(k, j) = \sum_{t \in T_{k,j}} E(t) / |T_{k,j}|$ .

We now create a regularized clustering cost function as

$$C_2(t) = \arg \max_{c=1..N} \left\{ \frac{e^{-\|f(t)-g(c)\|}}{\sum_{c=1}^N e^{-\|f(t)-g(c)\|}} + \lambda \frac{e^{-\frac{\|g(c)-g(C_2(t-1))\|}{E(t)}}}{\sum_{c=1}^N e^{-\frac{\|g(c)-g(C_2(t-1))\|}{E(c, C_2(t-1))}}} \right\} \quad (2.2)$$

where  $\lambda$  is the weighting parameter. Here motion energy  $E(t)$  plays a role as the temperature  $T$  in simulated annealing. When it is high (strong motion between frames), the motion continuity condition is violated and the labels of successive frames can change freely; when it is low, the smoothness term constrains the possible transitions of classes with low  $\bar{M}(k, j)$ .

With this in place, we now scan through the sequence searching for  $C_2(t)$  of maximum value given  $C_2(t-1)$  is already fixed.<sup>4</sup> This temporal smoothing is most relevant with images with motions, and static frames are already stably clustered and therefore their cluster labels do not change.

### GMM for Density Modeling and Smoothing

Given clusters, we build a probability density model for each. A Gaussian Mixture Model [43, 5] is used to gain good local relaxation based on the initial clustering result provided by the above method and good generalization for new data. Due to the curse of dimensionality, it is difficult to obtain a good estimate of a high dimensional density function with limited and largely varied training data. We introduce an iterative method incorporating Linear Discriminant Analysis (LDA) [5] and a GMM in an EM-like fashion to perform dimensional reduction. The initial clustering labels help to build the scatter matrices for LDA. The optimal projection matrix of LDA is then obtained from the decomposition of clusters' scatter matrices [5]. The original feature vectors can be further projected into a low dimensional space, which improves the estimation of multi-variate Gaussian density function.

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<sup>3</sup>A simple method is to compute motion energy as the Sum of Squared Differences (SSD) by subtracting two  $P_{skin}$  density masses from successive images.

<sup>4</sup>Note that  $\bar{M}(k, j)$  changes after scanning the labels of the image sequence once, thus more iterations could be used to achieve more accurate temporal smoothness of  $C_3(t)$ ,  $t = 1..N$ . From our experiments, more iterations does not change the result much.

With the new clustering result from GMM, LDA's scatter matrices and projection matrix can be re-estimated, and GMM can also be re-modeled in the new LDA subspace. This loop converges within  $5 \sim 10$  iterations from our experiments. Intuitively, LDA projects the data into a low dimensional subspace where the image clusters are well separated, which helps to have a good parameter estimation for GMM with limited data. Given more accurate GMM, more accurate clustering results are obtained, which also causes better estimate of LDA. This Discriminative-GMM algorithm is described in the following and the mathematical details are provided in the appendix. The algorithm includes 2 loops of iterations, the outer loop of LDA-GMM and the inner loop of GMM's EM process. Its experimental convergency is provided in section 3.5, in terms of the log-likelihood and misclassification rate of both real and synthesized data. The theoretical proof is currently undertaken. After this process, we have a Gaussian density model for each cluster.

### 2.2.3 High Level: Aggregated Histogram Model for Action Recognition

Given a set of  $n$  clusters, define  $w(t) = [p_{c_1}(f(t)), p_{c_2}(f(t)), \dots, p_{c_n}(f(t))]^T$  where  $p_x(y)$  denotes the density value of the vector  $y$  with respect to the GMM for cluster  $x$ . An action is then a trajectory of  $[w(t_1), w(t_1 + 1), \dots, w(t_2)]^T$  in  $\mathbb{R}^n$ . For recognition purposes, we want to calculate some discriminative statistics from each trajectory. One natural way is to use its mean  $H_{t_1, t_2} = \sum_{t=t_1}^{t_2} w(t)/(t_2 - t_1 + 1)$  over time which is a temporal weighted histogram. Note that the histogram  $H_{t_1, t_2}$  bins are precisely corresponding to the trained clusters.

From the training set, we aggregate the cluster weights of images within a given hand action to form a histogram model. In this way, a temporal image sequence corresponding to one action is represented by a single vector. The matching of different actions is equivalent to compute the similarity of two histograms which has variants. Here we use Bhattacharyya similarity metric [33] which has several useful properties including: it is an approximation of  $\chi^2$  test statistics with fixed bias; it is self-consistent; it does not have the singularity problem while matching empty histogram bins; and its value is properly bounded within  $[0, 1]$ . Assume we have a library of action histograms  $H_1^*, H_2^*, \dots, H_M^*$ , the class label of a new action  $\hat{H}_{t_1, t_2}$  is

### Discriminative-GMM Algorithm

*inputs:* Image Data  $\mathcal{X}$ ; Labels  $\mathcal{L}$ ; Mixture Number

$\mathcal{N}$

*outputs:* LDA Transform  $\mathcal{A}$ ; GMM Parameters

$\{\pi, \mu, \Sigma\}$

1. Construct  $\{\mathcal{S}_{\mathcal{W}}; \mathcal{S}_{\mathcal{B}}\} \Leftarrow \{\mathcal{X}; \mathcal{L}\}$ .
2. Compute LDA Transform  $\mathcal{A} \Leftarrow LDA(\mathcal{S}_{\mathcal{W}}; \mathcal{S}_{\mathcal{B}})$ , and project image data  $\mathcal{X}$  into the subspace  $\mathcal{Z} = \mathcal{A}^T \mathcal{X}$  with a lower dimensions  $\mathcal{D}$ .
3. Initialize GMM Parameters  $\{\pi, \mu, \Sigma\}$ :  $\pi_i = 1/\mathcal{N}$ ,  $\Sigma_i$  is a  $\mathcal{D}$  dimensional identical matrix where  $i = 1, 2, \dots, \mathcal{N}$ , and  $\mu_i \Leftarrow \{\mathcal{Z}; \mathcal{L}\}$  for the first time, while  $\mu_i \Leftarrow \{\mathcal{Z}; \mathcal{W}\}$  for others.
4. Learn GMM Parameters  $\{\pi, \mu, \Sigma\}$  for  $\mathcal{Z}$  via Expectation-Maximization iterations.
  - (a) E-step:  $\mathcal{W} \Leftarrow \{\mathcal{Z}; \{\pi, \mu, \Sigma\}\}$
  - (b) M-step:  $\{\pi, \mu, \Sigma\} \Leftarrow \{\mathcal{Z}; \mathcal{W}\}$
5. Construct  $\{\mathcal{S}_{\mathcal{W}}; \mathcal{S}_{\mathcal{B}}\} \Leftarrow \{\mathcal{X}; \mathcal{W}\}$ .
6. Go to (2), until convergence.

determined by the following equation.

$$L(\hat{H}_{t_1, t_2}) = \arg \min_{l=1..M} \{ D(H_l^*, \hat{H}_{t_1, t_2}) = \left[ 1 - \frac{1}{3} \sum_{c=1}^n \sqrt{H_l^*(c) * \hat{H}_{t_1, t_2}(c)} \right]^{\frac{1}{2}} \} \quad (2.3)$$

This method is low cost because only one exemplar per action category is needed.

One problem with this method is that all sequence information has been compressed, e.g., we cannot distinguish an opening hand gesture from a closing hand using only one histogram. This problem can be easily solved by subdividing the sequence and histogram model into  $m$  parts:  $H_{t_1,t_2}^m = [H_{t_1,(t_1+t_2)/m}, \dots, H_{(t_1+t_2)*(m-1)/m,t_2}]$ . For an extreme case when one frame is a subsequence, the histogram model simply becomes exactly the vector form of the representative surface.

We intend to classify hand actions with speed differences into the same category. To achieve this, the image frames within a hand action can be sub-sampled to build a set of temporal pyramids. In order to segment hand gestures from a long video sequence, we create several sliding windows with different frame sampling rates. The proper time scaling magnitude is found by searching for the best fit over temporal pyramids.

Taken together, the histogram representation achieves an adjustable multi-resolution measurement to describe actions. A Hidden Markov Model (HMM) with discrete observations could be also employed to train models for different hand actions, but more template samples per gesture class are required. The histogram recognition method has the additional advantage that it does not depend on extremely accurate frame-wise clustering. A small proportion of incorrect labels does not effect the matching value much. In comparison, in an HMM with few training samples, outliers seriously impact the accuracy of learning. From the viewpoint of considering hand actions as a language process, our model is an integration of individual observations (by labelling each frame with a set of learned clusters) from different time slots. The labels' transitions between successive frames are not used to describe the temporal sequence. By subdividing the histogram, we are extending the representation to contain bigram, trigram, etc. information.

## 2.3 Results

### 2.3.1 Convergency of Discriminative-GMM.

We first show the convergency of our Discriminative-GMM algorithm by the increasing log-likelihood curve of image data with the outer LDA-GMM iterations. In model selection literatures [5, 43], the model’s log-likelihood is a monotonic function of the number of clusters  $\mathcal{M}$ . In order to increase the log-likelihood given the learnt model, we need to increase  $\mathcal{M}$ . However we observe that the log-likelihood does increase significantly within the first several LDA-GMM loops while both the mixture model size and the subspace dimensions are fixed. We also generate 9 clusters of synthesized data with 5, 10, 15, 20, 25 dimensions. The first 2 dimensions are shown in Figure 2.4 (b), and other dimensions are not distinguishable because they are randomly sampled from the same normal distribution. Therefore, the optimal LDA projection should be a 2 dimensional subspace close to the first 2 dimensions. Figure 2.4 (c,d) illustrate the convergency of the LDA-GMM loops in terms of the improved log-likelihood and incorrect clustering ratios.

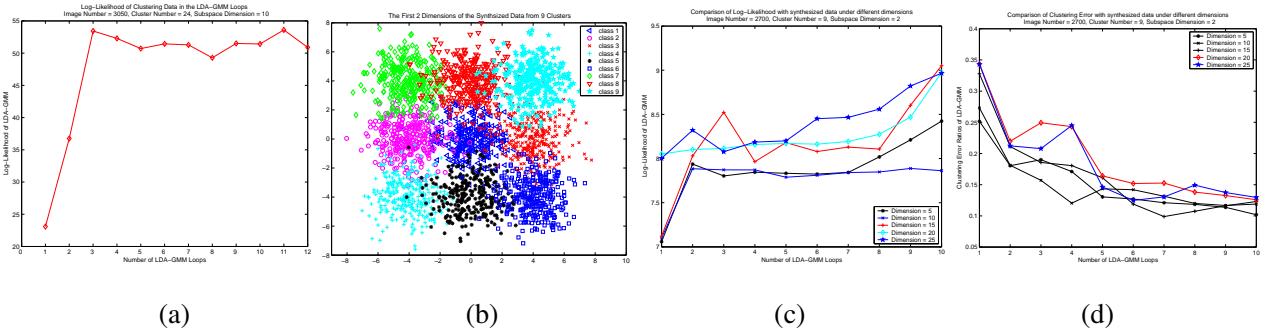


Figure 2.4: (a) Log-likelihood of 3015 images with 24 clusters and 10 dimensional subspace (b) The first 2 dimensions of the synthesized data from 9 clusters (c) Log-likelihood of the synthesized data of different dimensions (d) Ratios of incorrect clustering of the synthesized data of different dimensions.

We have tested our three tiered method on the problem of recognizing sequences of hand spelling gestures.

### 2.3.2 Framewise clustering.

We evaluate the low level representation of single images and intermediate clustering algorithms. A training set of 3015 images are used. The frame-to-frame motion energy is used to label images as static or dynamic.

2.0675e-006	2.3102e-018	1.6192e-010	6.0587e-013	5.7133e-009	4.5282e-009	2.0684e-013	1.1612e-002	8.2762e-002	2.1693e-001
Cluster 1	Cluster 3	Cluster 19	Cluster 5	Cluster 13	Cluster 8	Cluster 21	Cluster 15	Cluster 6	Cluster 12

Figure 2.5: Image clustering results after low and intermediate level processing.

For spectral clustering,  $3 \sim 4$  restarts from both the dynamic and static set are sufficient to cover all the modes in the training set. Then, temporal smoothing is employed and a Gaussian density is calculated for each cluster in a 10 dimensional subspace of the LDA projection. As a result, 24 clusters are obtained which contain 16 static and 8 dynamic modes. Figure 2.5 shows 5 frames closest to the mean of the probability density of cluster 1, 3, 19, 5, 13, 8, 21, 15, 6, 12. It can be seen that clustering results are insensitive to artifacts of skin segmentation. From Figure 2.5, it is also clear that dynamic modes have significantly larger determinants than static ones. The study of the eigenvalues of covariance matrices shows that their super-ellipsoid shapes are expanded within  $2 \sim 3$  dimensions or  $6 \sim 8$  dimensions for static or dynamic clusters. Taken together, this means that static clusters are quite tight, while dynamic clusters contain much more in-class variation. From Figure 2.6 (c), dynamic clusters gain more weight during the smoothing process incorporating the temporal constraint and subsequent GMM refinement.

### 2.3.3 Action recognition and segmentation.

For testing images, we first project their feature vectors into the LDA subspace. Then, the GMM is used to compute their weights with respect to each cluster. We manually choose 100 sequences for testing purposes, and compute their similarities with respect to a library of 25 gestures. The length of the action sequences

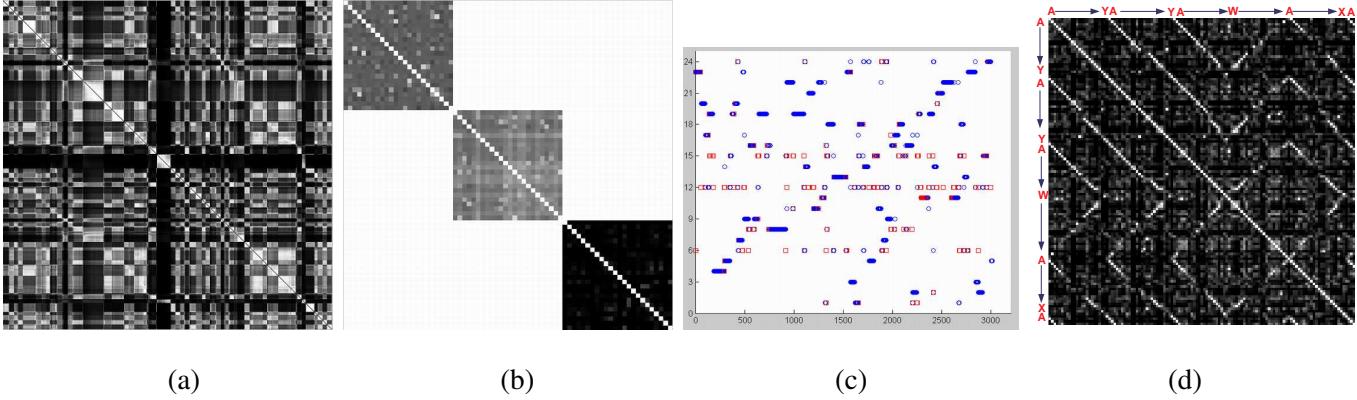


Figure 2.6: (a) Affinity matrix of 3015 images. (b) Affinity matrices of cluster centroids (from upper left to lower right) after spectral clustering, temporal smoothing and GMM. (c) Labelling results of 3015 images (red squares are frames whose labels changed with smoothing process after spectral clustering). (d) The similarity matrix of segmented hand gestures. The letters are labels of gestures, for example,  $A \rightarrow Y$  represents a sequence of gestures  $A \rightarrow B, B \rightarrow C, \dots, X \rightarrow Y$ .

was  $9 \sim 38$  frames. The temporal scale of actions in the same category ranged from 1 to 2.4. The results were recognition rates of 90% and 93% without/with temporal smoothing (Equation 2.2). Including the top three candidates, the recognition rates increase to 94% and 96%, respectively. We also used the learned model and a sliding window with temporal scaling to segment actions from a 6034 frame video sequence containing dynamic gestures and static hand postures. The similarity matrices among 123 actions found in the video is shown in Figure 2.6 (d). 106 out of 123 actions (86.2%) are correctly segmented and recognized.

### 2.3.4 Integrating motion information.

As noted previously, our method cannot distinguish opening/closing hand gestures without temporally subdividing histograms. An alternative solution is to integrate motion information<sup>5</sup> between frames. Motion feature vectors are also clustered, which results a joint (appearance and motion) histogram model for actions. We assume independence of the data and therefore simple contatenate these two histograms into a single action representation. From our preliminary experiments, both motion integration and histogram subdivision are comparably effective to recognize gestures with opposite direction.

<sup>5</sup>Motion information can be extracted by first aligning two hand blobs, subtracting two skin-color density masses, then using the same circular histogram in section 2.1 to extract a feature vector for positive and negative density residues respectively. Another simple way is to subtract two frames' feature vectors directly.

## 2.4 Conclusion and Discussion

We have presented a method for classifying the motion of articulated gestures using LDA/GMM-based clustering methods and a histogram-based model of temporal evolution. Using this model, we have obtained extremely good recognition results using a relatively coarse representation of appearance and motion in images.

There are mainly three methods to improve the performance of histogram-based classification, i.e., adaptive binning, adaptive subregion, and adaptive weighting [54]. In our approach, adaptive binning of the histogram is automatically learned by our clustering algorithms; adaptive subregion is realized by subdividing action sequences to enrich the histogram's descriptive capacity in the temporal domain; adaptive weighting is achieved from the trained weights of Gaussian kernels in GMM.

Our future work will focus on building a larger hand action database containing  $50 \sim 100$  categories for more extensive testing, and on extending the representation to include other types of image information (e.g. contour information). Also, by finding an effective foreground segmentation module, we intend to apply the same methods to other applications such as recognizing stylized human body motion.

## Chapter 3

# Discriminative Learning of Spatial Image

## Distribution:

# A Two-level Image Spatial Representation for Image Scene Category Recognition

*Classifying pictures into one of several semantic categories is a classical image understanding problem.*

*In this paper, we present a stratified approach to both binary (outdoor-indoor) and multiple category of scene classification. We first learn mixture models for 20 basic classes of local image content based on color and texture information. Once trained, these models are applied to a test image, and produce 20 probability density response maps (PDRM) indicating the likelihood that each image region was produced by each class. We then extract some very simple features from those PDRMs, and use them to train a bagged random Linear Discriminant Analysis (LDA) classifier for 10 scene categories. For this process, no explicit region segmentation or spatial context model are computed. To test this classification system, we created a labeled database of 1500 home photos taken under very different environment and lighting conditions, using different cameras, and from 43 persons over 5 years (1999-2004). The classification rate of outdoor-*

*indoor classification is 93.8%, and the classification rate for 10 scene categories is 90.1%. As a byproduct, local image patches can be contextually labeled into the 20 basic material classes by using Loopy Belief Propagation [167] as an anisotropic filter on PDRMs, producing an image-level segmentation if desired.*

### 3.1 Introduction

Classifying pictures into semantic types of scenes [151, 158, 144] is a classical image understanding problem which requires the effective interaction of high level semantic information and low level image observations. Our goal is to build a very practical prototype for scene classification of typical consumer photos, along the lines of the Kodak system [144]. Thus, we are interested in systems that are accurate, efficient, and which can work with a wide range of photos and photographic quality.

Given the extremely large within-category variations in typical photographs, it is usually simpler and thus easier to break the problem of scene classification into a two-step process. In this paper, we first train local, image patch based color-texture Gaussian Mixture models (GMM) to detect each of 20 materials in a local image patch. These models are used to scan an image and generate 20 local responses for each pixel. Each response map, called a Probability Density Response Map (PDRM), can be taken as a real-valued image indicating the relative likelihood of each material at each image location. We then compute moments from the response maps and form a feature vector for each photo. By employing the random subspace method [103, 162] and bootstrapping [165], we obtain a set of LDA scene classifiers over these feature vectors. These classification results are combined into the final decision through bagging [68]. After learning the local and global models, a typical  $1200 \times 800$  image can be classified in less than 1 second with our unoptimized Matlab implementation. Therefore there is a potential to develop a real-time scene classifier upon our approach. A complete diagram of our approach is shown in Figure 3.1.

We organize the rest of the paper as follows. After a brief review of related previous published work, we present the local image-level processing used to create PDRMs In section 3.3. Then in 3.4, we describe how PDRMs are processed to perform scene classification. Experimental results and analysis on the performance of patch based material detector and image based scene classification on a database of 1500 personal photos

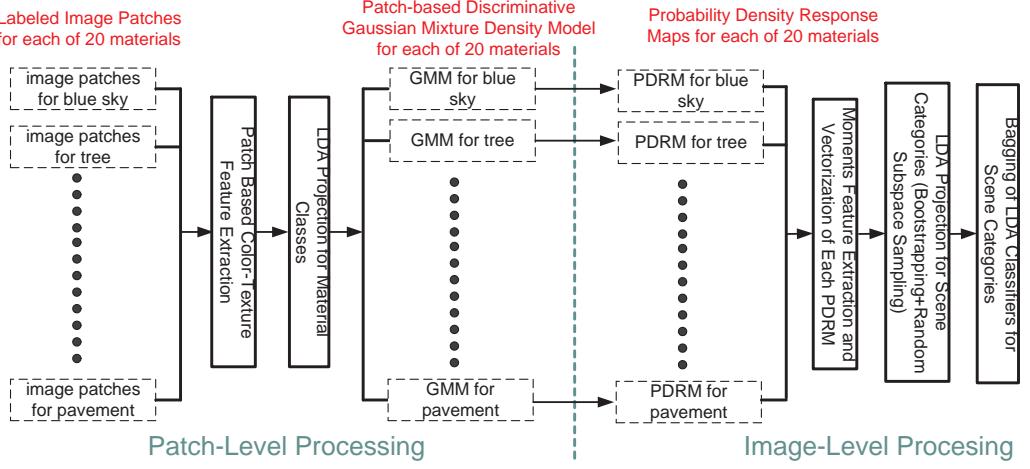


Figure 3.1: The diagram of our two level approach for scene recognition. The dashed line boxes are the input data or output learned models; the solid line boxes represent the functions of our algorithm.

taken by 43 users using traditional or digital cameras over the last 5 years are given in section 3.5. Finally we summarize the paper and discuss the future work in section 3.6.

## 3.2 Previous Work

There are several related efforts in this area. Luo et al. [127, 144] propose a bottom-up approach to first find and label well-segmented image regions, such as water, beach, sky, and then to learn the spatial contextual model among regions. A Bayesian network codes these relational dependencies. By comparison, we do not perform an explicit spatial segmentation, and we use relatively simple (LDA-based) classification methods. Perona et al. [83, 164] present a constellation model of clustered feature components for object recognition. Their method works well for detecting single objects, but strongly depends on the performance and reliability of the interest detector [110]. In the case of scene classification, we need to model more than one class of material, where classes are non-structural and do not have significant features (such as foliage, rock and et al.) [110]. This motivates our use of a GMM on the feature space. In order to maintain good stability, we estimate the GMM in a linear subspace computed by LDA. These density models are quite flexible and can be used to model a wide variety of image patterns with a good compromise between discrimination and smoothness.

Kumar et al. [114, 115] propose the use of Markov random field (MRF)-based spatial contextual models to detect man-made buildings in a natural landscape. They build a multi-scale color and textual descriptor to capture the local dependence among building and non-building image blocks and use MRF to model the prior of block labels. In our work, we have found that simple local labeling suffices to generate good classification results; indeed regularization using loopy belief propagation method [167] yields no significant improvement in performance. Thus, we claim that there is no need to segment image regions explicitly for scene classification as other authors have done [144, 127, 115].

Linear discriminant analysis (LDA) is an optimization method to compute linear combinations of features that have more power to separate different classes. For texture modeling, Zhu et al [169] pursue features to find the marginal distributions which are also the linear combinations of the basic filter banks, but they use a much more complex method (Monte Carlo Markov Chain) to stochastically search the space of linear coefficients. In our case, the goal is not to build a generative model for photos belonging to different scenes, but simply to discriminate among them. We show a simple method such as LDA, if designed properly, can be very effective and efficient to build a useful classifier for complex scenes.

### 3.3 Local Image-Level Processing

The role of image-level processing is to roughly classify local image content at each location in the image. The general approach is to compute feature vectors of both color and texture, and then develop classifiers for these features. In our current implementation, we have chosen to perform supervised feature classification. Although arguably less practical than corresponding unsupervised methods, supervised classification permits us to control the structure of the representations built at this level, and thereby to better understand the relationship between low-level representations and overall system performance.

In this step, we compute 20 data driven probabilistic density models to describe the color-texture properties of image patches of 20 predefined materials<sup>1</sup>. These 20 categories are: building, blue sky, bush, other

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<sup>1</sup>The vocabulary of materials to be detected is designed by considering their popularity in the usual family photos. This definition is, of course, not unique or optimized.

(mostly trained with human clothes), cloudy sky, dirt, mammal, pavement, pebble, rock, sand, skin, tree, water, shining sky, grass, snow, carpet, wall and furniture.

To prepare the training data, we manually crop image regions for each material in our database, and randomly draw dozens of 25 by 25 pixel patches from each rectangle. Altogether, we have 2000 image patches for each material. Some examples of the cropped images and sampled image patches are shown in Figure 3.2. For simplicity, we do not precisely follow the material boundaries in the photos while cropping. Some outlier features are thus included in the training patches. Fortunately these outliers are smoothed nicely by learning continuous mixture density models.

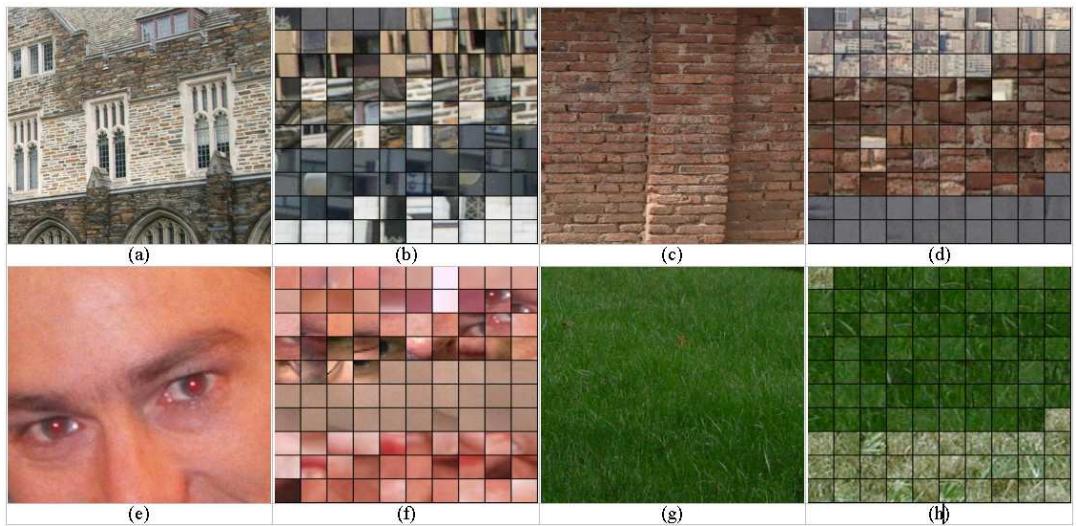


Figure 3.2: (a, c, e, g) Examples of cropped subimages of building, building under closer view, human skin, and grass respectively. (b, d, f, h) Examples of image patches of these materials including local patches sampled from the above subimages. Each local image patch is 25 by 25 pixels.

Multi-scale image representation and automatic scale selection problem has been a topic of intense discussion over the last decade [125, 128, 110, 78, 114]. In general, the approach of most authors has been to first normalize images with respect to the estimated scale of local image regions before learning. However it is not a trivial problem to reliably recover the local image scales for a collection of 1500 family photos. We instead choose to train the GMM using the raw image patches extracted directly from the original pictures. For the labeled image patches with closer and coarser views, their complex color-texture distributions can

will be approximated by a multi-modal Gaussian mixture model during clustering.

### 3.3.1 Color-Texture Descriptor for Image Patches

Our first problem is to extract a good color-texture descriptor which effectively allows us to distinguish the appearance of different materials. In the domain of color, experimental evaluation of several color models has not indicated significant performance differences among color representations. As a result, we simply represent the color of an image patch as the mean color in RGB space.

There are also several methods to extract texture feature vectors for image patches. Here we consider two: filter banks, and the Haralick texture descriptor. Filter banks have been widely used for 2 and 3 dimensional texture recognition. [121, 76, 160]. We apply the **Leung-Malik (LM) filter bank** [121] which consists of 48 isotropic and anisotropic filters with 6 directions, 3 scales and 2 phases. Thus, each patch is represented by a 48 component feature vector.

The Haralick texture descriptor [98] is designed for image classification and has been adopted in the area of image retrieval [61]. Haralick texture measurements are derived from the Gray Level Co-occurrence Matrix (GLCM). GLCM is also called the **Grey Tone Spatial Dependency Matrix** which is a tabulation of how often different combinations of pixel brightness values (grey levels) occur in an image region. GLCM texture considers the relation between two pixels at a time, called the reference and the neighbor pixel. Their spatial relation can be decided by two factors, the orientation and offset. Given any image patch, we search all the pixel pairs satisfying a certain spatial relation and record their second order gray level distributions with a 2 dimensional histogram indexed by their brightness values<sup>2</sup>. Haralick also designed 14 different texture features [98] based on the GLCM. We selected 5 texture features including dissimilarity, Angular Second Moment (ASM), mean, standard deviation (STD) and correction. Definitions for these can be found in Appendix 6.1.

There is no general argument that the filter bank features or Haralick feature is a better texture descriptor. We evaluate their texture discrimination performances experimentally in section 3.5 and find Haralick

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<sup>2</sup>The reference and neighbor pixel intensities normally need to be quantized into 16 or less levels instead of 256 which results in not too sparse GLCM.

features generally perform better.

### 3.3.2 Discriminative Mixture Density Models for 20 Materials

The color and texture features for 2000 image patches form, in principle, an empirical model for each material. However, classifying new patches against the raw features would require the solution to a high-dimensional nearest-neighbor problem, and the result would be sensitive to noise and outliers. Instead, we compute a continuous membership function using a Gaussian mixture model.

Although we have 2000 training samples, our feature vectors have 40 dimensions, so the training set is still too sparse to learn a good mixture model without dimensional reduction. Because one of our purposes is to maximize the discrimination among different materials, Linear Discriminant Analysis (LDA) [165] was chosen to project the data into a subspace where each class is well separated. The LDA computation is reviewed in appendix 6.2.

When each class has a Gaussian density with a common covariance matrix, LDA is the optimal transform to separate data from different classes. Unfortunately the material color-texture distributions all have multiple modes because the training image patches are sampled from a large variety of photos. Therefore we have two options: employ LDA to discriminate among 20 material classes; or use LDA to separate all the modes of materials. Although the latter seems closer to the model for which LDA was designed, we found its material classification rate is worse because the optimal separation among the multiple modes within the same material class is irrelevant. Therefore we choose the former.

The LDA computation provides a projection of the original feature space into a lower-dimensional feature space  $\mathcal{Z}$ . We assume that the color-texture features of each material class is described by a finite mixture distribution on  $\mathcal{Z}$  of the form

$$P(z|c) = \sum_{k=1}^{g_c} \pi_k^c \mathcal{G}(z; \mu_k^c, \Sigma_k^c), \quad c = 1, 2, \dots, 20 \quad (3.1)$$

where the  $\pi_k^c$  are the mixing proportions ( $\sum_{k=1}^{g_c} \pi_k^c = 1$ ) and  $\mathcal{G}(z; \mu_k^c, \Sigma_k^c)$  is a multivariate Gaussian function depending on a parameter vector  $\theta_k^c$ . The number of mixtures  $g_c$  and the model parameters  $\{\pi_k^c, \theta_k^c\}$  for each material class  $c$  are initialized by spectral clustering [135] and learned in an iterative Expectation-

Maximization manner [165, 40] where  $g_c$  ranged from 4 to 8 depending on the material class. As a summary, discriminative Gaussian mixture models are obtained by applying LDA across the material classes and learning the GMM within each material class, respectively.

## 3.4 Global Image Processing

### 3.4.1 Global Image Descriptor

Once we obtain 20 Gaussian mixture models  $\{\pi_k^i, P(z; \theta_k^i), i = 1, 2, \dots, 20\}$  for 20 material classes, we can evaluate the membership density values of image patches for each material class. For any given photo, we scan local image patches, extract their color-texture feature vector, normalize each of its components from 0 to 1 [61], project it to the lower dimensional subspace  $\mathcal{Z}$  computed by LDA, and finally compute the density value given by equation (3.1) for all 20 material classes. The result is 20 real-valued grid maps<sup>3</sup> representing membership support for each of the 20 classes. An example is shown in Figure 3.3. Two examples of the local patch labeling for indoor and outdoor photos are shown in Figure 3.4.

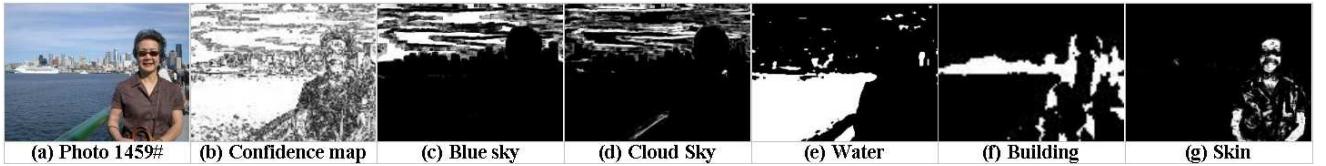


Figure 3.3: (a) Photo 1459#. (b) Its confidence map. (c, d, e, f, g) Its support maps of blue sky, cloud sky, water, building and skin. Only the material classes with the significant membership support are shown.

Our next goal is to classify the photos into one of ten categories: cityscape, landscape, mountain, beach, snow, other outdoors, portrait, party, still life and other indoor. In order to classify photos, we must still reduce the dimension of the PDRMs to a manageable size. To do this, we compute the zeroth, first, and second order moments of each PDRM. Intuitively, the zeroth moment describes the prevalence of a given material class in an image; the first moment describes where it occurs, and the second moment its spatial "spread". The moment features from the 20 PDRMs are combined in a global feature vector  $Y$ .

<sup>3</sup>The size of the map depends on the original photo size and the patches' spatial sampling intervals.



(a)

(b)

Figure 3.4: (a) The local patch material labeling results of an indoor photo. (b) The local patch material labeling results of an outdoor photo. Loopy belief propagation is used for enhancement. The colored dots represent the material label and the boundaries are manually overlayed for illustration purpose only.

### 3.4.2 Week Learner: LDA Classifiers with Bootstrapping and Random Subspace Sampling

Using the scene category labels of the training photos, we now compute the LDA transform that attempts to separate the training feature vectors of different categories. For the indoor-outdoor recognition, the LDA projected subspace has only one dimension. As a typical pattern classification problem, we can find the optimal decision boundary from the training data and apply it to the other testing data. Finding decision boundaries for 10 scene category recognition is more complex. In practice, it is very difficult to train a GMM classifier because of the data is too sparse over the 10 categories. As a result, we have used both the nearest neighbor and Kmeans [165] classifiers for this decision problem.

We have found that the standard method for creating an LDA classifier works well for indoor-outdoor scene classification, but the classification results for 10 scene categories is not good enough to constitute a practical prototype. To improve the classification rate, we have implemented variations on random subspace generation [103, 162] and bootstrapping [165] to create multiple LDA classifiers. These classifiers are combined using bagging [68]. Recall that LDA is a two step process that first computes the singular value decomposition (SVD) [90] of the within-class scatter matrix  $\mathbf{S}_W$ , then, after normalization, computes SVD on the between-class scatter matrix  $\mathbf{S}'_B$ . After the first step,  $\mathbf{S}_W$  is divided into the principal subspace  $\mathbf{S}_P$  of the nonzero eigenvalues  $\Lambda_P$  and their associated eigenvectors  $\mathbf{U}_P$ , and the null subspace  $\mathbf{S}_N$  with the zero eigenvalues  $\Lambda_N$  and corresponding eigenvectors  $\mathbf{U}_N$ . In the traditional LDA transform, only  $\mathbf{S}_P$  is used for the whitening of  $\mathbf{S}_W$  and normalization of  $\mathbf{S}_B$  while  $\mathbf{S}_N$  is discarded (see equation 6.18 in Appendix 6.2).

Chen et al. [72] have found that the null subspace  $\mathbf{S}_N$  satisfying  $\mathbf{U}_P^T \mathbf{S}_W \mathbf{U}_P = 0$  also contains important discriminatory information. Here we make use of this observation by uniformly sampling an eigenvector matrix  $\mathbf{U}_r$  from  $\{\mathbf{U}_P \cup \mathbf{U}_N\}$  and use it in place of  $\mathbf{U}$  in the initial LDA projection step. Several projections (including the original LDA projection matrix) are thus created.

In the second step of LDA, the subset  $\mathbf{V}_P$  of the full eigenvector matrix  $\mathbf{V}$  with the largest eigenvalues, normally replaces  $\mathbf{V}$  in equation (6.18). It is also possible that there is useful discriminative information in the subspace  $\{\mathbf{V} - \mathbf{V}_P\}$ . Therefore we employ a similar sampling strategy as [162] in the context of PCA by first sampling a small subset of eigenvectors  $\mathbf{V}_r$  of  $\{\mathbf{V} - \mathbf{V}_P\}$ , then replacing  $\mathbf{V}$  with the joint subspace  $\{\mathbf{V}_P \cup \mathbf{V}_r\}$  in equation 6.18.

Finally we also perform bootstrapping [165] by sampling subjects of the training set and creating LDA classifiers for these subsets. By the above three random sampling processes, we learn a large set of LDA subspaces and classifiers which we combine using the majority voting (bagging) methods [68]. In Section 3.5, we show the bagged recognition rates of 20 classifiers from bootstrapping replicates and 20 from random subspace sampling.

## 3.5 Experiments

Our photo collection currently consists of 540 indoor and 860 outdoor customer photos. We randomly select half of them as the training data and use other photos as the testing data. We have also intentionally minimized redundancy when collecting photos, i.e., only one photo is selected when there are several similar pictures.

### 3.5.1 Local Recognition: Validation of Image Patches Representation for Material Classes

We first address the problem of the image patch based color-texture feature description and classification. Comparison of the recognition rates of 1200 testing image patches for each material class for different color-texture descriptors, different numbers of training patches and different classifiers is provided in Figure 3.6 (a,b). In particular, we have also benchmarked the LDA+GMM model against a brute-force nearest neighbor

classifier. Let  $x_j$  and  $z_j$  represent an image patch feature vector before and after the LDA projection, respectively. The nearest neighbor classifier computes the class label of a testing patch  $j$  as the label of that training patch  $l$  such that  $\|x_j - x_l\| = \min_i \{\|x_j - x_i\|\}$  where  $i$  ranges over the training image patches of all material classes. The GMM classifier simply chooses the maximal class density, i.e. the class  $c^*$  such that  $P(z_j|c^*) = \max_{c=1,2,\dots,20} \{P(z_j|c)\}$ .

Comparing the plots shown in Figure 3.6, the classifier based on the Maximum Likelihood of GMM density functions outperforms the Nearest Neighbor classifier, thus validating the use of the LDA+GMM method. We also compared the recognition rates of 4 different feature combinations and found that the Haralick texture descriptor combined with the mean color of the image patch yields the best results. Finally, in Figure 3.6 (b), we see that the LDA+GMM method improves the recognition rate significantly when increasing the training image patch from 500, becoming stable after 2000 patches.

Figure 3.5 shows the confusion rate using the GMM classifiers learned from 2000 training image patches per class. The size of the white rectangle in each grid is proportional to the pairwise recognition error ratio. The largest and smallest confusion rates are 23.6% and 0.24%, respectively. From Figure 3.5, we see that pebble, rock and sand classes are well separated which shows that our patch-level learning process achieves a good balance of Haralick texture and color cues by finding differences of the material classes with the similar color. There is significant confusion among grass, bush and tree due to their similar color and texture distribution. For some material classes, such as furniture, carpet, and other, the overall confusion rates are also high.

### 3.5.2 Global Recognition: Photo based Scene Category Classification

For global classification, we have found that first order moment features of PRDMs are useful in outdoor scenes, but reduce the recognition rate for indoor scenes. This makes sense since in most outdoor scenes spatial contextual constraints, for instance the sky above grass, are useful cues. This naturally suggests a hierarchical classification scheme (first determine indoor/outdoor followed by categorization), however we have not yet pursued this approach. Thus, we confine ourselves to zeroth order moments for the remainder

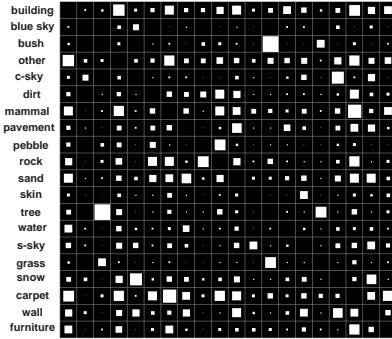


Figure 3.5: The pairwise confusion matrix of 20 material classes. The indexing order of the confusion matrix is shown on the left of the matrix. The indexing order is symmetrical.

of this paper.

Our global image moment features after LDA projection are very easy to visualize in the indoor/outdoor case as they become points in a 1-dimensional LDA subspace (3.6 (c)). In this case, the 1-D indoor-outdoor decision boundary is simply determined by fitting a scaled exponential function to each of the indoor or outdoor histogram distributions and calculating the point of intersection.

We show the recognition results of our method in Figure 3.6 (d), compared with the direct low-level color or texture based scene recognition methods<sup>4</sup> without LDA learning as the baselines. Our indoor-outdoor recognition rate is 93.8%, which is comparable or slightly better than the Kodak's recently published classification system [144], although our approach is tested on a 40% larger photo database. It is interesting that the bagging algorithm does not significantly improve the recognition performance of for indoor-outdoor classification. The likely explanation is that the individual indoor-outdoor LDA classifiers have nearly achieved the best possible recognition rate. Figure 3.7 shows 2 examples of misclassified photos. The first photo consists of a person sitting indoors, but in front of a curtain of tree leaves. In the second, the playground is incorrectly classified as "carpet" not "dirt". The appearance of people and animals are irrelevant for indoor-outdoor classification — their associated moment features are assigned with near zero

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<sup>4</sup>We divide each image as a 9 by 9 grid, and extract the mean color or the DOG (Derivative of Gaussian) filtered texture features within each grid. Each photo is then formulated as a feature vector by combining cues in all grids. A nearest neighbor classifier is later employed for recognition based on the feature vectors' distances of the training and testing photos.

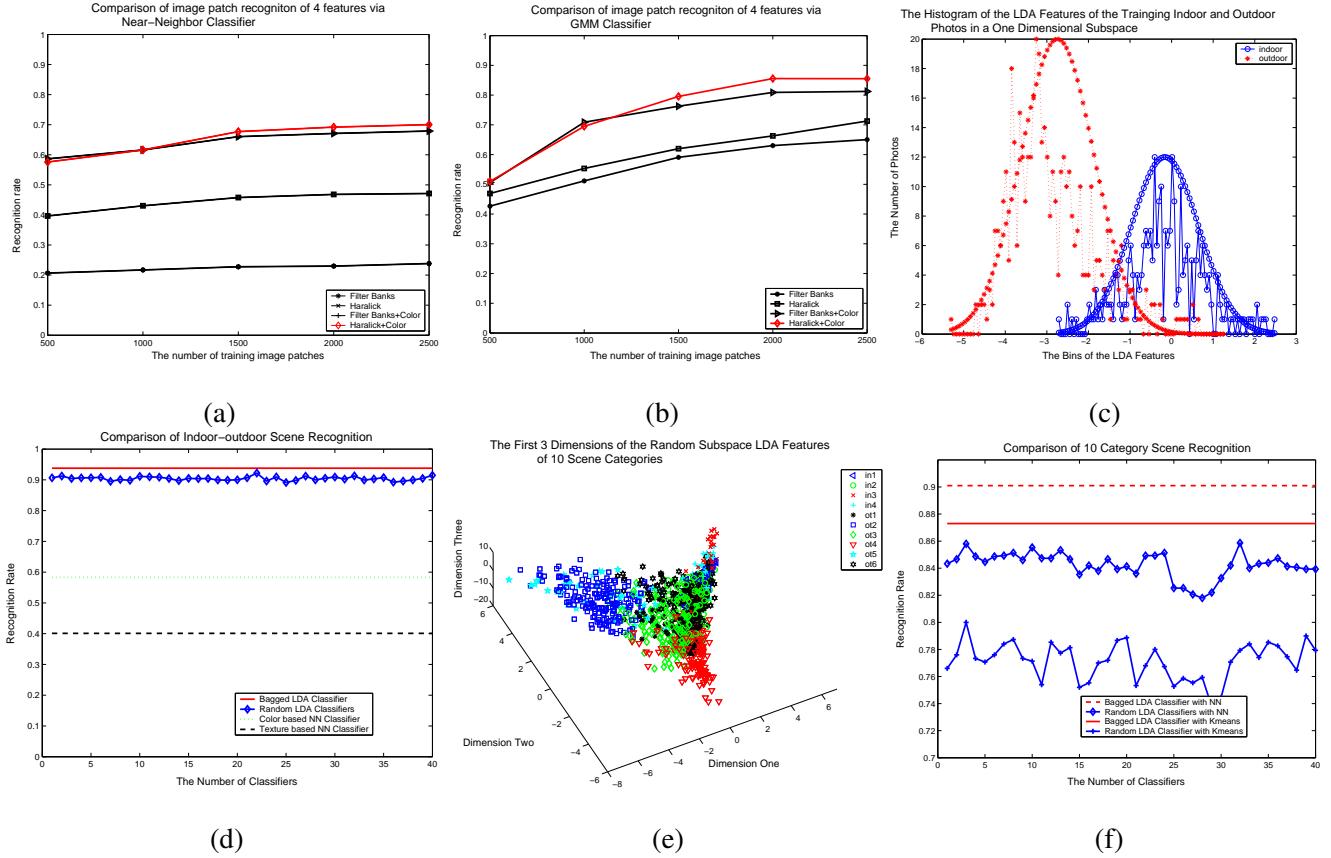


Figure 3.6: (a) Comparison of the image patch based recognition of 4 kinds of features (filter banks feature, Haralick texture feature and their joint features with color) via Nearest-Neighbor Classifier. (b) Comparison of the image patch based recognition of 4 kinds of features via GMM Classifier. (c)The 1D feature histogram distributions of indoor-outdoor photos after LDA projection. (d) The comparison of indoor-outdoor recognition rates of 4 methods. (e) The first 3D feature point distributions of 10 category photos after LDA projection. (f) The comparison of 10 categories recognition rates of 4 methods.

weights.

As shown in Figure 3.6 (e), feature points of some scene categories are well separated from others and thus easy to be recognized in a certain LDA subspace, while some categories are not. Fortunately, Figure 3.6 (f) demonstrates that the individual LDA classifiers capture the complimentary discriminative information in different random subspaces. Finally, it results that the combined (nearest neighbor and Kmeans) classifiers both show improved performances of 6 – 10% on average. As a comparison, Boutell et al. [66] achieve less than 80% classification accuracy for 923 images in 5 categories. In their work, model-based graph matching techniques are used to learn the explicit scene configuration consisting of semantic image regions.



(a)



(b)

Figure 3.7: (a) An misclassified indoor photo. (b) An misclassified outdoor photo.

### 3.6 Conclusions & Discussion

This paper makes three contributions. First, we propose an efficient, yet effective, approach for scene recognition for both indoor-outdoor and multiple photo categories. In practice, this approach can handle the photos' spatial complexity both in the local patch-level and the global image-level successfully. All the training and testing processes are based upon a challenging photo database. Second, we describe a combination of LDA and Gaussian mixture models that achieves a good balance of discrimination and smoothness. Finally, we study the use of moment features of PDRMs as an effective image-level representation for scene classification, and the bagging [68] method to combine the individual scene classifiers obtained by the random subspace algorithm [103]. The bagging method has shown success in our experiments, especially for 10 category scene recognition.

Although we have used supervised methods to create the local image patch classifiers, a practical system would like learn at least some of these classifiers using unsupervised methods. However we believe that the supervised material detectors provide the best scene recognition performance, and as such provide a “benchmark” against which unsupervised methods can be evaluated. In future work, we intend to investigate unsupervised clustering methods for low-level image patch classification. In particular, we plan to apply our unsupervised, iterative LDA-GMM algorithm [?]. We also plan to investigate a hybrid approach where classified images are used as labeled data to compute an initial LDA projection, which is then subsequently refined with new, unlabeled images using iterative LDA-GMM. Finally, because LDA is only optimal when each class has a Gaussian density with a common covariance matrix, the non-parametric discriminant analysis (proposed in [168]) will be tested as a means to generalize our approach to a more comprehensive image

database which may contain thousands of various kinds of photos.

## **Chapter 4**

### **Online Learning of Dynamic**

### **Spatial-Temporal Image and Video**

#### **Appearance:**

#### **Matching, Segmenting and Tracking**

#### **Objects in Images and Videos using**

#### **Non-parametric Random Image Patch**

#### **Propagation**

*In this paper, we propose a novel exemplar-based approach to extract dynamic foreground regions from a changing background within a collection of images or a video sequence. By using image segmentation as a pre-processing step, we convert this traditional pixel-wise labeling problem into a lower-dimensional supervised, binary labeling procedure on image segments. Our approach consists of three steps. First, a set*

*of random image patches are spatially and adaptively sampled within each segment. Second, these sets of extracted samples are formed into two “bags of patches” to model the foreground/background appearance, respectively. We perform a novel bidirectional consistency check between new patches from incoming frames and current “bags of patches” to reject outliers, control model rigidity and make the model adaptive to new observations. Within each bag, image patches are further partitioned and resampled to create an evolving appearance model. Finally, the foreground/background decision over segments in an image is formulated using an aggregation function defined on the similarity measurements of sampled patches relative to the foreground and background models. The essence of the algorithm is conceptually simple and can be easily implemented within a few hundreds of lines of Matlab code. We evaluate and validate the proposed approach by extensive real examples of the object-level image mapping and tracking within a variety of challenging environments. We also show that it is straightforward to apply our problem formulation on non-rigid object tracking with difficult surveillance videos.*

## 4.1 Introduction

In this paper, we study the problem of object-level figure/ground segmentation in images and video sequences. The core problem can be defined as follows: Given an image  $\mathbb{X}$  with known figure/ground labels  $\mathbb{L}$ , infer the figure/ground labels  $\mathbb{L}'$  of a new image  $\mathbb{X}'$  closely related to  $\mathbb{X}$ . For example, we may want to extract a walking person in an image using the figure/ground mask of the same person in another image of the same sequence. Our approach is based on training a classifier from the appearance of a pixel and its surrounding context (i.e., an image patch centered at the pixel) to recognize other similar pixels across images. To apply this process to a video sequence, we also evolve the appearance model over time.

A key element of our approach is the use of a prior segmentation to reduce the complexity of the segmentation process. As argued in [138], image segments are a more natural primitive for image modeling than pixels. More specifically, an image segmentation provides a natural dimensional reduction from the spatial resolution of the image to a much smaller set of spatially compact and relatively homogeneous regions. We can then focus on representing the appearance characteristics of these regions. Borrowing a term

from [138], we can think of each region as a "superpixel" which represents a complex connected spatial region of the image using a rich set of derived image features. We can then consider how to classify each superpixel (i.e. image segment) as foreground or background, and then project this classification back into the original image to create the pixel-level foreground-background segmentation we are interested in.

The original superpixel representation in [138, 129, 128] is a feature vector created from the image segment's color histogram [129], filter bank responses [138], oriented energy [128] and contourness [128]. These features are effective for image segmentation [128], or finding perceptually important boundaries from segmentation by supervised training [138]. However, as shown in [11], those parameters do not work well for matching different classes of image regions from different images. Instead, we propose using a set of spatially randomly sampled image patches as a non-parametric, statistical superpixel representation. This non-parametric "bag of patches" model<sup>1</sup> can be easily and robustly evolved with the spatial-temporal appearance information from video, while maintaining the model size (the number of image patches per bag) using adaptive sampling. Foreground/background classification is then posed as the problem of matching sets of random patches from the image with these models. Our *major contributions* are demonstrating the effectiveness and computational simplicity of a nonparametric random patch representation for semantically labelling superpixels and a novel bidirectional consistency check and resampling strategy for robust foreground/background appearance adaptation over time.

We organize the paper as follows. We first address the related work in the areas of computer vision and graphics. Then several image patch based representations and the associated matching methods are described. In section 4.4, the algorithm used in our approach is presented with details. We demonstrate the validity of the proposed approach using experiments on real examples of the object-level figure/ground image mapping and non-rigid object tracking under dynamic conditions from videos of different resolutions in section 4.5. Finally, we summarize the contributions of the paper and discuss possible extensions and improvements.

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<sup>1</sup>Highly distinctive local features [126] are not the adequate substitutes for image patches. Their spatial sparseness nature limits their representativity within each individual image segment, especially for the nonrigid, nonstructural and flexible foreground/background appearance.

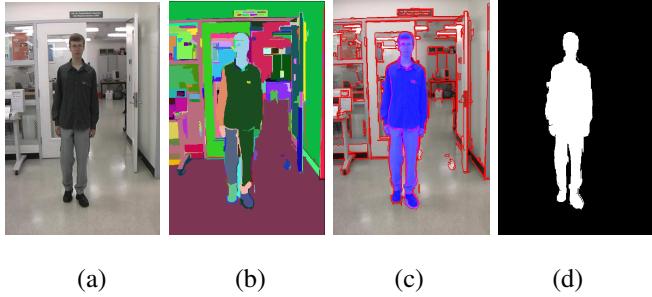


Figure 4.1: (a) An example indoor image, (b) the segmentation result using [82] coded in random colors, (c) the boundary pixels between segments shown in red, the image segments associated with the foreground, a walking person here, shown in blue, (d) the associated foreground/background mask. Notice that the color in (a) is not very saturated. This is a common fact in our indoor experiments without any specific lighting controls.

## 4.2 Related work

In this paper, we propose a simple nonparametric random patch treatment unifying supervised semantic object segmentation across images, non-rigid object segmentation tracking in moderate/high resolution videos and temporal small object localization in low resolution videos. All these tasks involve a non-rigid figure object, a moving camera or a changing background. We show real examples in a variety of conditions which are related to a wide range of previous work as follows.

Most previous work on foreground/background extraction are based on the pixel-level processing of images from static cameras using color as the primary feature [73, 139, 123, 124, 161, 133, 146]. Image patch based appearance modeling [80, 121, 11] and matching enriches both the descriptive and discriminative abilities for figure/ground classification compared to the pixel-color representation. Patch based image sampling and matching has shown great success in texture synthesis [80], texture recognition [121] and superior performance in object recognition [?] than image features [126].

Interactively extracting a foreground object from an image [139, 123], or segmenting a moving object from a video sequence [124, 161] remains a difficult computer graphics task. The state-of-the-art [139, 123, 124, 161] employ an interactive graph-cut algorithm [67] as a Markov random field solver to assign pixels with figure/ground labels using color cues. It normally needs huge amount of manual interactions and assumes the camera pose is simply fixed. Our paper attempts to provide an automatic means to propagate

supervised semantic segmentation labels over images by nonparametric appearance modeling and temporal-spatial video frames.

Li et al. [123] and Rother et al. [139] utilized an interactive graph-cut algorithm [67] as a Markov random field solver to assign pixels with figure/ground labels. Li et al. [124] further extended this approach to video cutout applications using a 3D graph-cut algorithm on the spatial-temporal space. Most work is primarily based on color, and video cutout papers [124, 161] assume that the background is static.

Dynamically changing backgrounds render many of the above methods ineffective. In recent work, [146, 133] describe pixel-wise foreground detection algorithms to handle a quasi-static<sup>2</sup> background. This work relies on a local smoothing process on the pixels occupied by dynamic textures using a kernel density estimator in the joint spatial-color space. However, the approach does not handle the change in background due to a moving camera. Motion segmentation is another approach to find independent moving objects by computing an exact model of background motion [?]. Unfortunately it is only effective for segmenting small moving objects from a constrained dominant background motion, mostly for aerial visual surveillance applications. By comparison, our treatment of image segments (instead of pixels) as the elements of foreground/background classification avoids the need for motion assumptions across images.

The idea of using "superpixels" as the representative elements for object-level image mapping is inspired by [138, 129, 123, 105]. For moderate resolution images, the number of segments in a segmentation is typically several orders of magnitude lower than the number of pixels. This makes the foreground/background extraction problem computationally more tractable. Also, as we will show, moderately rigid or non-rigid spatial transforms of figure/ground across images tend not to affect our segment-based classification. As a result, there is no reliance on a figure/ground motion or shape model. Our non-parametric statistical appearance representation of superpixel is in the same spirit of [80]. They employ it to synthesize textures, while we use it for segment-based foreground/background labelling. Recently, [?] proposes a hierarchical model switching method for unsupervised video segmentation which involves variational inference over many conditional switching and conditional hidden variables. It is very computationally expensive (thus feasible for

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<sup>2</sup>A static scene with periodically changing objects, such as a running river, waving trees, or ocean waves and so on.

limited low resolution videos only), and depends highly on a complex switching process among different global shape/appearance models.

We also extend our nonparametric formulation of random patches into tracking non-rigid interested objects from surveillance videos, and compared it with the state-of-the-art [?, 96]. [96] utilizes meanshift mode seeking [?] to maintain an online appearance-changing Gaussian mixture density model. The meanshift density model has potential difficulties for high dimensional image features that limit its descriptiveness. [?] employs Adaboosting to obtain an ensemble of recently trained weak classifiers for the temporal appearance model updating and classification. This approach loses the original format of image patch appearance and restricts to boosting for classification. Instead of using boosting as in [?], our approach has a novel bidirectional check and resampling strategy to keep the figure/ground dynamic appearance model robust and updated along time. As we show later, the strategy only involves simple computations with flexible and controllable descriptive ability while naturally combining long-term and short-term appearance samples. It also leaves the classification process with freedom to choose over different classifiers.

### 4.3 Image Patch Representation and Matching

Building stable appearance representations of images patches is fundamental to our approach. There are many derived features that can be used to represent the appearance of an image patch. In this paper, we evaluate our algorithm based on: 1) an image patch's raw RGB intensity vector, 2) mean color vector, 3) color + texture descriptor (filter bank response or Haralick feature [11]), and 4) PCA, LDA and NDA (Nonparametric Discriminant Analysis) features [87, 69] on the raw RGB vectors. For completeness, we give a brief description of each of these techniques.

#### 4.3.1 Texture descriptors

To compute texture descriptions, we first apply the *Leung-Malik (LM) filter bank* [121] which consists of 48 isotropic and anisotropic filters with 6 directions, 3 scales and 2 phases. Thus each image patch is represented by a 48 component feature vector. The *Haralick texture descriptor* [98] was used for image

classification in [11]. Haralick features are derived from the Gray Level Co-occurrence Matrix, which is a tabulation of how often different combinations of pixel brightness values (grey levels) occur in an image region. We selected 5 out of 14 texture descriptors [98] including dissimilarity, Angular Second Moment (ASM), mean, standard deviation (STD) and correlation. For details, refer to [98, 11].

### 4.3.2 Dimension reduction representations

The *Principal Component Analysis* (PCA) algorithm is used to reduce the dimensionality of the raw color intensity vectors of image patches. PCA makes no prior assumptions about the labels of data. However, recall that we construct the "bag of patches" appearance model from sets of labelled image patches. This supervised information can be used to project the bags of patches into a subspace where they are best separated using *Linear discriminant Analysis* (LDA) or *Nonparametric Discriminant Analysis* (NDA) algorithm [87, 69] by assuming Gaussian or Non-Gaussian class-specific distributions.

#### Comparison of LDA and NDA

Both NDA and LDA compute an optimal projection matrix  $W$  based on the within-class and between-class scatter matrices  $S_W, S_B$ <sup>3</sup>, by maximizing the intra-class separation  $\{W^T S_B W / W^T S_W W\}$ . When each class has a Gaussian density with a common covariance matrix, LDA is the optimal discriminative transform to separate data from different classes. However the image patches of foreground/background classes usually have very complex multimodal distributions. The nonparametric nature of scatter matrices  $S_W, S_B$  in NDA [87, 69] can inherently lead to extract projected features that preserve relevant complex structures of classification. However, through our experiments, the nonlinear separability of different classes in the NDA projected subspace is very sensitive to its parameter settings while building scatter matrices. There is no principled way or computationally intractable to find optimal parameters for arbitrary complex distributions.

NDA differs from LDA in how it constructs  $S_W, S_B$  matrices. For each image patch  $p$ , we need to find

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<sup>3</sup> $S_W$  is the covariance matrix of data to its intra-class mean;  $S_B$  is the covariance matrix of the intra-class means to the overall mean.

the means  $\bar{p}^I, \bar{p}^E$  of its nearest neighbor sets  $\{p^I\}, \{p^E\}$  from both the intra-class and inter-class patch bags. This can be computationally expensive with the large size bags of patches and the high dimensionality of the image patch. In this paper, we cluster image patches within each bag (as described in section 4.4.4) and use the cluster centers to find approximations of  $\bar{p}^I, \bar{p}^E$  as follows. Given the center sets  $C^F, C^B$ , any foreground image patch's intra-class mean is chosen as  $c \in C^F$  within the same partition and its inter-class mean is  $c \in C^B$  with the minimal distance; similarly for background patches. Then  $S_W, S_B$  are constructed as covariance matrices using these local means [69]. The computational complexity decreases from  $O(N^2d)$  to  $O(kNd)$  where  $N$  image patches are clustered into  $k$  partitions and  $d$  is the patch feature vector's dimensionality. After constructing the parametric or non-parametric scatter matrices  $S_W, S_B$ , both LDA and NDA can be solved as a generalized eigenvalue problem [5]. For details, refer to [87, 69].

### 4.3.3 Patch matching

: After image patches are represented using one of the above methods, we must match them against the foreground/background models. There are 2 methods investigated in this paper: the nearest neighbor matching using Euclidean distance and KDE (Kernel Density Estimation) [106] in PCA/NDA subspaces. For nearest-neighbor matching, we find, for each patch  $p$ , its nearest neighbors  $p_n^F, p_n^B$  in foreground/background bags, and then compute  $d_p^F = \|p - p_n^F\|$ ,  $d_p^B = \|p - p_n^B\|$ . On the other hand, an image patch's matching scores  $m_p^F$  and  $m_p^B$  are evaluated as probability density values from the KDE functions  $KDE(p, \Omega^F)$  and  $KDE(p, \Omega^B)$  where  $\Omega^{F|B}$  are bags of patch models. Then the segmentation-level classification is performed as section 4.4.3.

## 4.4 Algorithms

We briefly summarize our labeling algorithm as follows. We assume that each image of interest has been segmented into spatial regions. A set of random image patches are spatially and adaptively sampled within each segment. These sets of extracted samples are formed into two “bags of patches” to model the foreground/background appearance respectively. The foreground/background decision for any segment in a new

image is computed using one of two aggregation functions on the appearance similarities from its inside image patches to the foreground and background models. Finally, for videos, within each bag, new patches from new frames are integrated through a robust bidirectional consistency check process and all image patches are then partitioned and resampled to create an evolving appearance model. As described below, this process prune classification inaccuracies in the nonparametric image patch representations and adapts them towards current changes in foreground/background appearances for videos.

#### 4.4.1 Algorithm Diagram

We describe each of these steps for video tracking of foreground/background segments in more detail below, and for image matching, which we treat as a special case by simply omitting step 3 and 4 in Figure 4.2.

#### 4.4.2 Sample Random Image Patches

We first employ an image segmentation algorithm<sup>4</sup> [82] to pre-segment all the images or video frames in our experiments. A typical segmentation result is shown in Figure 4.1. We use  $\mathbb{X}_t, t = 1, 2, \dots, T$  to represent a sequence of video frames. Given an image segment, we formulate its representation as a distribution on the appearance variation over all possible extracted image patches inside the segment. To keep this representation to a manageable size, we approximate this distribution by sampling a random subset of patches.

We denote an image segment as  $\mathcal{S}_i$  with  $\mathcal{S}_i^F$  for a foreground segment, and  $\mathcal{S}_i^B$  for a background segment, where  $i$  is the index of the (foreground/background)image segment within an image. Accordingly,  $\mathcal{P}_i, \mathcal{P}_i^F$  and  $\mathcal{P}_i^B$  represent a set of random image patches sampled from  $\mathcal{S}_i, \mathcal{S}_i^F$  and  $\mathcal{S}_i^B$  respectively. The cardinality  $\mathcal{N}_i$  of an image segment  $\mathcal{S}_i$  generated by [82] typically ranges from 50 to thousands. However small or large superpixels are expected to have roughly the same amount of uniformity. Therefore the sampling rate  $\gamma_i$  of  $\mathcal{S}_i$ , defined as  $\gamma_i = \text{size}(\mathcal{P}_i)/\mathcal{N}_i$ , should decrease with increasing  $\mathcal{N}_i$ . For simplicity, we keep  $\gamma_i$  as a constant for all superpixels, unless  $\mathcal{N}_i$  is above a predefined threshold  $\tau$ , (typically  $2500 \sim 3000$ ), above

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<sup>4</sup>Because we are not focused on image segmentation algorithms, we choose Felzenszwalb's segmentation code which generates good results and is publicly available at <http://people.cs.uchicago.edu/~pff/segment/>.

### Non-parametric Patch Appearance Modelling-Matching Algorithm

*inputs:* Pre-segmented Images  $\mathbb{X}_t, t = 1, 2, \dots, T$ ; Label  $\mathbb{L}_1$

*outputs:* Labels  $\mathbb{L}_t, t = 2, \dots, T$ ; 2 “bags of patches” appearance model for foreground/background  $\Omega_T^{F|B}$

1. Sample segmentation-adaptive random image patches  $\{\mathcal{P}_1\}$  from image  $\mathbb{X}_1$ .
2. Construct 2 new bags of patches  $\Omega_1^{F|B}$  for foreground/background using patches  $\{\mathcal{P}_1\}$  and label  $\mathbb{L}_1$ ; set  $t = 1$ .
3.  $t = t + 1$ ; Sample segmentation-adaptive random image patches  $\{\mathcal{P}_t\}$  from image  $\mathbb{X}_t$ ; match  $\{\mathcal{P}_t\}$  with  $\Omega_{t-1}^{F|B}$  and classify segments of  $\mathbb{X}_t$  to generate label  $\mathbb{L}_t$  by aggregation.
4. Classify and reject ambiguous patch samples, probable outliers and redundant appearance patch samples from new extracted image patches  $\{\mathcal{P}_t\}$  against  $\Omega_{t-1}^{F|B}$ ; Then integrate the filtered  $\{\mathcal{P}_t\}$  into  $\Omega_{t-1}^{F|B}$  and evaluate the probability of survival  $p_s$  for each patch inside  $\Omega_{t-1}^{F|B}$  against the original unprocessed  $\{\mathcal{P}_t\}$  (Bidirectional Consistency Check).
5. Perform the random partition and resampling process according to the normalized product of probability of survival  $p_s$  and partition-wise sampling rate  $\gamma'$  inside  $\Omega_{t-1}^{F|B}$  to generate  $\Omega_t^{F|B}$ .
6. If  $t = T$ , output  $\mathbb{L}_t, t = 2, \dots, T$  and  $\Omega_T^{F|B}$ ; exit. If  $t < T$ , go to (3).

Figure 4.2: Non-parametric Patch Appearance Modelling-Matching Algorithm

which  $\text{size}(\mathcal{P}_i)$  is held fixed. This sampling adaptivity is illustrated in Figure 4.3. Notice that large image segments have much more sparsely sampled patches than small image segments. From our experiments, this adaptive spatial sampling strategy is sufficient to represent image segments of different sizes.

#### 4.4.3 Label Segments by Aggregating Over Random Patches

For an image segment  $\mathcal{S}_i$  from a new frame to be classified, we again first sample a set of random patches  $\mathcal{P}_i$  as its representative set of appearance samples. For each patch  $p \in \mathcal{P}_i$ , we calculate its distances  $d_p^F, d_p^B$  or matching scores  $m_p^B, m_p^F$  towards the foreground and background appearance models respectively as



Figure 4.3: **Left:** Segment adaptive random patch sampling from an image with known figure/ground labels. Green dots are samples for background; dark brown dots are samples for foreground. **Right:** Segment adaptive random patch sampling from a new image for figure/ground classification, shown as blue dots.

described in Section 4.3.

The decision of assigning  $\mathcal{S}_i$  to foreground or background, is an aggregating process over all  $\{d_p^F, d_p^B\}$  or  $\{m_p^B; m_p^F\}$  where  $p \in \mathcal{P}_i$ . Since  $\mathcal{P}_i$  is considered as a set of i.i.d. samples of the appearance distribution of  $\mathcal{S}_i$ , we use the average of  $\{d_p^F, d_p^B\}$  or  $\{m_p^B; m_p^F\}$  (ie. first-order statistics) as its distances  $D_{\mathcal{P}_i}^F, D_{\mathcal{P}_i}^B$  or fitness values  $M_{\mathcal{P}_i}^F, M_{\mathcal{P}_i}^B$  with the foreground/background model. In terms of distances  $\{d_p^F, d_p^B\}$ ,  $D_{\mathcal{P}_i}^F = \text{mean}_{p \in \mathcal{P}_i}(d_p^F)$  and  $D_{\mathcal{P}_i}^B = \text{mean}_{p \in \mathcal{P}_i}(d_p^B)$ . Then the segment's foreground/background fitness is set as the inverse of the distances:  $M_{\mathcal{P}_i}^F = 1/D_{\mathcal{P}_i}^F$  and  $M_{\mathcal{P}_i}^B = 1/D_{\mathcal{P}_i}^B$ . In terms of KDE matching scores  $\{m_p^B; m_p^F\}$ ,  $M_{\mathcal{P}_i}^F = \text{mean}_{p \in \mathcal{P}_i}(m_p^F)$  and  $M_{\mathcal{P}_i}^B = \text{mean}_{p \in \mathcal{P}_i}(m_p^B)$ . Finally,  $\mathcal{S}_i$  is classified as foreground if  $M_{\mathcal{P}_i}^F > M_{\mathcal{P}_i}^B$ , and vice versa. The *Median* robust operator can also be employed in our experiments, without noticeable difference in performance. Another choice is to classify each  $p \in \mathcal{P}_i$  from  $m_p^B$  and  $m_p^F$ , then vote the majority foreground/background decision for  $\mathcal{S}_i$ . The performance is similar with *mean* and *median*.

#### 4.4.4 Construct a Robust Online Nonparametric Foreground/Background Appearance Model with Temporal Adaptation

From sets of random image patches extracted from superpixels with known figure/ground labels, 2 foreground/background ‘‘bags of patches’’ are composed. The bags are the non-parametric form of the foreground/background appearance distributions. When we intend to ‘‘track’’ the figure/ground model sequentially through a sequence, these models need to be updated by integrating new image patches extracted from new video frames. However the size (the number of patches) of the bag will be unacceptably large if we do not also remove the some redundant information over time. More importantly, imperfect segmentation results from [82] can cause inaccurate segmentation level figure/ground labels. For robust image patch level appearance modeling of  $\Omega_t$ , we propose a novel bidirectional consistency check and resampling strategy to tackle various noise and labelling uncertainties.

More precisely, we classify new extracted image patches  $\{\mathcal{P}_t\}$  as  $\{\mathcal{P}_t^F\}$  or  $\{\mathcal{P}_t^B\}$  according to  $\Omega_{t-1}^{F|B}$ ; and reject ambiguous patch samples whose distances  $d_p^F, d_p^B$  towards respective  $\Omega_{t-1}^{F|B}$  have no good contrast (simply, the ratio between  $d_p^F$  and  $d_p^B$  falls into the range of 0.8 to 1/0.8). We further sort the distance list of the newly classified foreground patches  $\{\mathcal{P}_t^F\}$  to  $\Omega_{t-1}^F$ , filter out image patches on the top of the list which have too large distances and are probably to be outliers, and ones from the bottom of the list which have too small distances and contain probably redundant appearances compared with  $\Omega_{t-1}^F$ <sup>5</sup>. We perform the same process with  $\{\mathcal{P}_t^B\}$  according to  $\Omega_{t-1}^B$ . Then the filtered  $\{\mathcal{P}_t\}$  are integrated into  $\Omega_{t-1}^{F|B}$  to form  $\Omega_{t-1}^{F'|B'}$ , and we evaluate the probability of survival  $p_s$  for each patch inside  $\Omega_{t-1}^{F'|B'}$  against the original unprocessed  $\{\mathcal{P}_t\}$  with their labels<sup>6</sup>.

Next, we cluster all image patches of  $\Omega_{t-1}^{F'|B'}$  into  $k$  partitions [91], and randomly resample image patches within each partition. This is roughly equivalent to finding the modes of an arbitrary distribution and sam-

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<sup>5</sup>Simply, we reject patches with distances  $d_{p_t^F}^F$  that are larger than  $\text{mean}(d_{p_t^F}^F) + \lambda * \text{std}(d_{p_t^F}^F)$  or smaller than  $\text{mean}(d_{p_t^F}^F) - \lambda * \text{std}(d_{p_t^F}^F)$  where  $\lambda$  controls the range of accepting patch samples of  $\Omega_{t-1}^{F|B}$ , called *model rigidity*.

<sup>6</sup>For example, we compute the distance of each patch in  $\Omega_{t-1}^{F'}$  to  $\{\mathcal{P}_t^F\}$ , and covert them as surviving probabilities using a exponential function over negative covariance normalized distances. Patches with smaller distances have higher survival chances during resampling; and vice versa. We perform the same process with  $\Omega_{t-1}^{B'}$  according to  $\{\mathcal{P}_t^B\}$ .

pling for each mode. Ideally, the resampling rate  $\gamma'$  should decrease with increasing partition size, similar to the segment-wise sampling rate  $\gamma$ . For simplicity, we define  $\gamma'$  as a constant value for all partitions, unless setting a threshold  $\tau'$  to be the minimal required size<sup>7</sup> of partitions after resampling. If we perform resampling directly over patches without partitioning, some modes of the appearance distribution may be mistakenly removed. This strategy represents all partitions with sufficient number of image patches, regardless of their different sizes. In all, we resample image patches of  $\Omega_{t-1}^{F|B}$ , according to the normalized product of probability of survival  $p_s$  and partition-wise sampling rate  $\gamma'$ , to generate  $\Omega_t^{F|B}$ . By approximately fixing the expected bag model size, the number of image patches extracted from a certain frame  $\mathbb{X}_t$  in the bag decays exponentially in time.

The problem of partitioning image patches in the bag can be formulated as the NP-hard *k-center* problem. The definition of *k-center* is as follows: given a data set of  $n$  points and a predefined cluster number  $k$ , find a partition of the points into  $k$  subgroups  $\mathcal{P}_1, \mathcal{P}_2, \dots, \mathcal{P}_k$  and the data centers  $c_1, c_2, \dots, c_k$ , to minimize the maximum radius of clusters  $\max_i \max_{p \in \mathcal{P}_i} \| p - c_i \|$ , where  $i$  is the index of clusters. Gonzalez [91] proposed an efficient greedy algorithm, *farthest-point clustering*, which proved to give an approximation factor of 2 of the optimum. The algorithm operates as follows: pick a random point  $p_1$  as the first cluster center and add it to the center set  $C$ ; for iterations  $i = 2, \dots, k$ , find the point  $p_i$  with the farthest distance to the current center set  $C$ :  $d_i(p_i, C) = \min_{c \in C} \| p_i - c \|$  and add  $p_i$  to set  $C$ ; finally assign data points to its nearest center and recompute the means of clusters in  $C$ . Compared with the popular k-means algorithm, this algorithm is computationally efficient and theoretically bounded<sup>8</sup>. In this paper, we employ the Euclidean distance between an image patch and a cluster center, using the raw RGB intensity vector or the feature representations discussed in section 4.3.

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<sup>7</sup>All image patches from partitions that are already smaller than  $\tau'$  are kept during resampling.

<sup>8</sup>The random initialization of all  $k$  centers and the local iterative smoothing process in k-means, which is time-consuming in high dimensional space and possibly converges to undesirable local minimum, are avoided.

## 4.5 Experiments

We have evaluated the image patch representations described in Section 4.3 for figure/ground mapping between pairs of image on video sequences taken with both static and moving cameras. Here we summarize our results.

### 4.5.1 Evaluation on Object-level Figure/Ground Image Mapping

We first evaluate our algorithm on object-level figure/ground mapping between pairs of images under eight configurations of different image patch representations and matching criteria. They are listed as follows: the nearest neighbor distance matching on the image patch's mean color vector (*MCV*); raw color intensity vector of regular patch scanning (*RCV*) or segment-adaptive patch sampling over image (*SCV*); color + filter bank response (*CFB*); color + Haralick texture descriptor (*CHA*); PCA feature vector (*PCA*); NDA feature vector (*NDA*) and kernel density evaluation on PCA features (*KDE*). In general,  $8000 \sim 12000$  random patches are sampled per image. There is no apparent difference on classification accuracy for the patch size ranging from 9 to 15 pixels and the sample rate from 0.04 to 0.10. The PCA/NDA feature vector has 20 dimensions, and KDE is evaluated on the first 3 PCA features.

Because the foreground figure has fewer of pixels than background, we conservatively measure the classification accuracy from the foreground's detection precision and recall on pixels. Precision is the ratio of the number of correctly detected foreground pixels to the total number of detected foreground pixels; recall is the ratio of the number of correctly detected foreground pixels to the total number of foreground pixels in the image.

The patch size is 11 by 11 pixels, and the segment-wise patch sampling rate  $\gamma$  is fixed as 0.06. Using 40 pairs of  $(720 \times 480)$  images with the labelled figure/ground segmentation, we compare their average running time and classification accuracies in Tables 4.1 and 4.2. All the algorithms are implemented under Matlab 6.5 on a P4-1.8G PC.

The nearest neighbor matching has computational complexity  $O(N^2d)$  where  $N$  is the number of sampled patches per image and  $d$  is the dimensionality of the image patch representation. Therefore the running

<i>MCV</i>	<i>RCV</i>	<i>SCV</i>	<i>CFB</i>	<i>CHA</i>	<i>PCA</i>	<i>NDA</i>	<i>KDE</i>
0.9	8.5	4.5	2.2	2.6	1.2	1.6	0.38

Table 4.1: Evaluation on running time (minutes).

<i>MCV</i>	<i>RCV</i>	<i>SCV</i>	<i>CFB</i>	<i>CHA</i>	<i>PCA</i>	<i>NDA</i>	<i>KDE</i>
0.46	0.81	0.97	0.92	0.89	0.93	0.96	0.69
0.28	0.89	0.95	0.85	0.81	0.85	0.87	0.98

Table 4.2: Evaluation on classification accuracy (ratio). The first row is precision; the second row is recall. time differences of *MCV*, *RCV*, *SCV*, *CFB*, *CHA*, *PCA*, *NDA* mostly depend on  $d$ , except for the extra expense of feature extraction for *CFB*, *CHA*, *PCA*, *NDA*. Given an 11 by 11 pixel image patch, its raw RGB intensity vector has 363 dimensions. The dimensionality of color-texture descriptor is 51 for *LM* filter bank and 43 for Haralick texture features. The PCA and NDA features have dimensionalities ranging from 5 to 40 with comparable classification accuracy. *KDE* Matlab toolbox [106] uses tree-based approximations on kernel evaluation which boosts its speed. For figure/ground extraction accuracy, *SCV* has the best classification ratio using the raw color intensity vector without any dimension reduction. *MCV* has the worst accuracy, which shows that pixel-color leads to poor separability between figure and ground in our data set. Four feature based representations, *CFB*, *CHA*, *PCA*, *NDA* with reduced dimensions, have similar performance, whereas *NDA* is slightly better than the others. *KDE* tends to be more biased towards the foreground class because background usually has a wider, flatter density distribution. The superiority of *SCV* over *RCV* proves that our segment-wise random patch sampling strategy is more effective at classifying image segments than regularly scanning the image, even with more samples. As shown in Figure 4.4 (b), some small or irregularly-shaped image segments do not have enough patch samples to produce stable classifications.

To allow a more intuitive understanding the evaluation results, we present a set of example images with the detected figure segments in blue and their white/black masks in Figure 4.4. Another example of figure/ground mapping results from surgical images is illustrated in Figure 4.6. Our approach successfully distinguishes the extracted foreground object from another surgical tool with very similar appearance in the

background. More examples of mapping various figure/ground objects from one template image to other testing images are shown in supplementary materials.

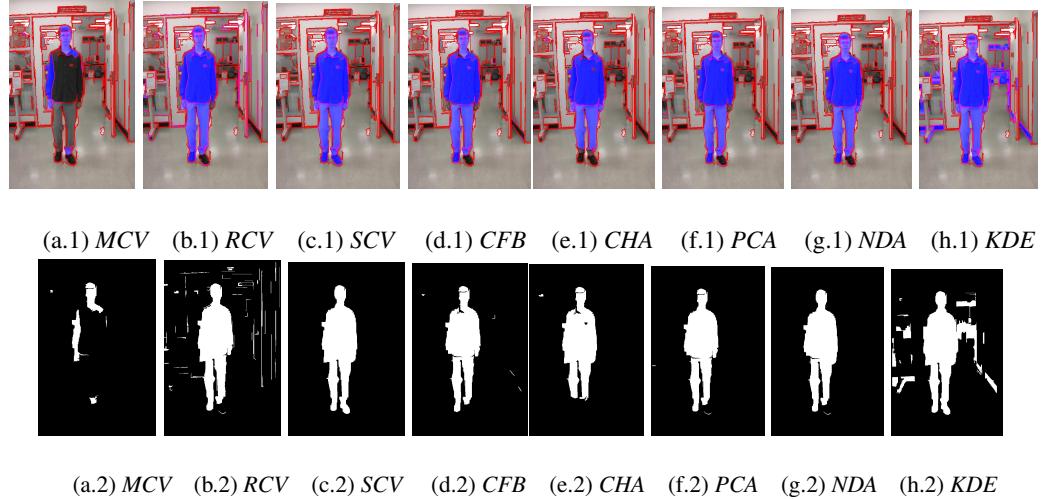
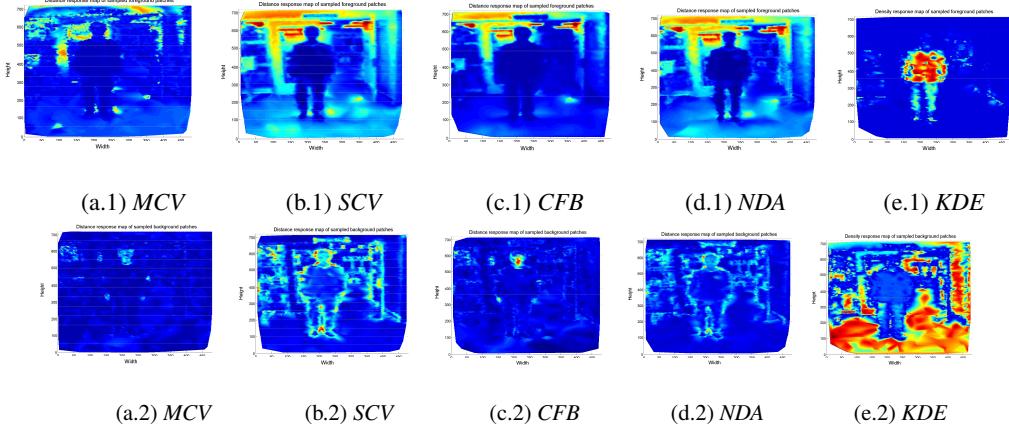


Figure 4.4: An example of evaluation on object-level figure/ground image mapping. The images with detected figure segments coded in blue are shown in the first row; their corresponding image masks are presented in the second row.

#### 4.5.2 Figure/Ground Segmentation Tracking with a Moving Camera

From Figure 4.4 (h), we see *KDE* tends to produce some false positives for the foreground. However the problem can be effectively tackled by multiplying the appearance *KDE* by the spatial prior which is also formulated as a *KDE* function of image patch coordinates. By considering videos with complex appearance-changing figure/ground, imperfect segmentation results [82] are not completely avoidable which can cause superpixel based figure/ground labelling errors. However our *robust bidirectional consistency check and resampling strategy*, as shown below, enables to successfully track the dynamic figure/ground segmentations in challenging scenarios with outlier rejection, model rigidity control and temporal adaptation (as described in section 4.4.4).

*Karsten.avi* shows a person walking in an uncontrolled indoor environment while tracked with a hand-held camera. After we manually label the frame 1, the foreground/background appearance model starts to develop, classify new frames and get updated online. Eight Example tracking frames are shown in Figure 4.7. Notice that the significant non-rigid deformations and large scale changes of the walking person, while



**Figure 4.5:** An example of the “bags of patches” model matching distance maps in (a,b,c,d) and density map in (e), within the image coordinates. Red means larger value; blue means smaller value. Smaller distances and larger density values represent better model-matching fitness, and vice versa. Due to space limits, we only show the results of *MCV*, *SCV*, *CFB*, *NDA*, *KDE* for the foreground model matching in the first row and background model matching in the second row. Compared to *SCV*, *CFB*, *NDA*, *RCV*, *CHA*, *PCA* have very similar distance maps.

the original background is completely substituted after the subject turned his way. In frame 258, we manually eliminate some false positives of the figure. The reason for this failure is that some image regions which were behind the subject begin to appear when the person is walking from left to the center of image (starting from frame 220). Compared to the online foreground/background appearance models by then, these newly appearing image regions have quite different appearance from both the foreground and the background. Thus the foreground’s spatial prior dominates the classification. There is another tracking example at an outdoor scenario in Figure 4.8.

**Automatic initialization:** As an extension of the basic methods, we also automatically detect the foreground from a handheld video camera. To do so, we first capture a few frames of the background without foreground appearance and extract random patches to fill into the background patch bag. Then foreground detection becomes an outlier detection problem and the newly detected foreground (outlier) segments are sampled into the foreground bag. Finally we iterate the foreground/background classification and the bag of patches model building process to convergence. This process depends on an outlier threshold setting and is sensitive to viewpoint changes during the capture of background vs. the initial foreground/background

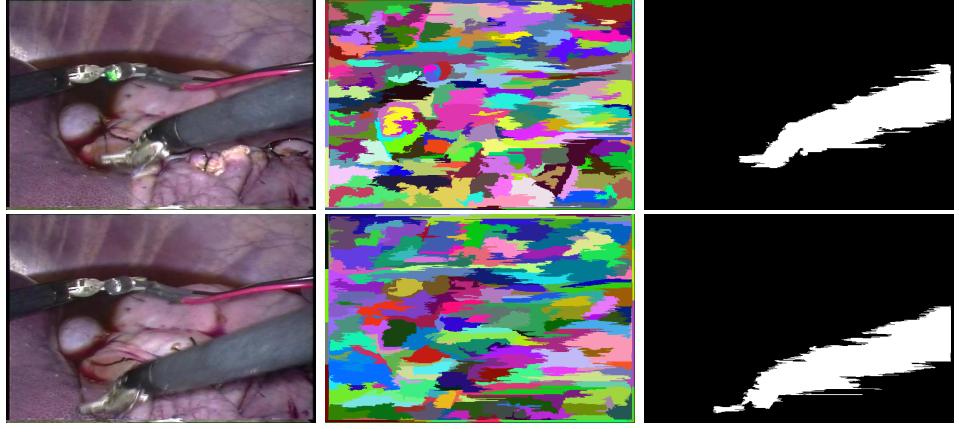


Figure 4.6: **Top Left:** An image for learning the foreground/background appearance model; **Top Middle:** Its segmentation; **Top Right:** Its labelling mask (White is foreground; black is background); **Bottom Left:** Another image for testing the appearance model; **Bottom Middle:** Its segmentation; **Bottom Right:** Its detected foreground/background mask. We use the patch based raw RGB intensity vector matching and the nearest neighbor matching. Notice the motions between 2 images. Image resolution is 720 by 488 pixels.

frame. Thus the iterations do not always converge to the desirable foreground/background separation. We show one of our successful results in Figure 4.9. Further improvements are under investigation.

#### 4.5.3 Non-rigid Object Tracking from Surveillance Videos

We can also apply our nonparametric treatment of dynamic random patches in Figure 4.2 into tracking non-rigid interested objects from surveillance videos. The difficult is that surveillance cameras normally capture small non-rigid figures, such as a walking person or running car, in low contrast and low resolution format. Thus to adapt our method to solve this problem, we make the following modifications. Because our task changes to localizing figure object automatically overtime, we can simply model figure/ground regions using rectangles (as shown in Figure 4.10) and therefore no pre-segmentation [82] is needed. Random figure/ground patches are then extracted from the image regions within these two rectangles. Using two sets of random image patches, we train an online classifier for figure/ground classes at each time step, generate a figure appearance confidence map of classification for the next frame and, similarly to [?], apply mean shift [?] to find the next object location by mode seeking. In our problem solution, the temporal evolution of dynamic image patch appearance models are executed by the bidirectional consistency check and resampling

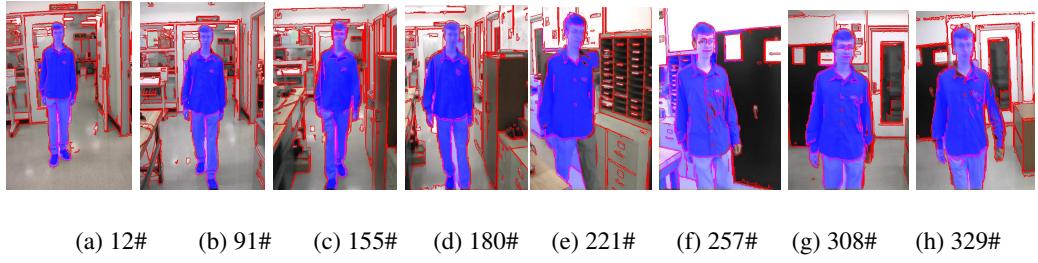


Figure 4.7: Eight example frames (720 by 480 pixels) from the video sequence *Karsten.avi* of 330

frames. The video is captured using a handheld Panasonic PV-GS120 in standard NTSC format. Notice that the significant non-rigid deformations and large scale changes of the walking person, while the original background is completely substituted after the subject turned his way. The red pixels are on the boundary of segments; the tracked image segments associated with the foreground walking person is coded in blue.

described in section 4.4.4. Whereas [?] uses boosting for both temporal appearance model updating and classification, our online binary classification training can employ any off-the-shelf classifiers. We have tested on Nearest Neighbors (NN), discriminative probability density models (DPDM) [11] and support vector machine (SVM). DPDM performs slightly worse than NN and SVM by considering the difficulty of building density functions over complexly distributed and temporally evolving appearance samples. In Figure 4.10, we demonstrate sample frames from four video sequences using SVM classification and our bidirectional consistency check and resampling technique. Our results are comparable with the state-of-the-art [?, 96], even under more challenging scenario such as Figure 4.10 (d). Our tracking method performs directly on the original video resolutions, instead of using three layers of image pyramids [?]. *Refer supplementary materials for more details.*

## 4.6 Conclusion and Discussion

Although quite simple both conceptually and computationally, our algorithm of performing dynamic foreground-background extraction in images and videos using non-parametric appearance models produces very promising and reliable results in a wide variety of circumstances. For tracking figure/ground segments, to our best knowledge, it is the first attempt to solve this difficult "video matting" problem [124, 161] by robust and

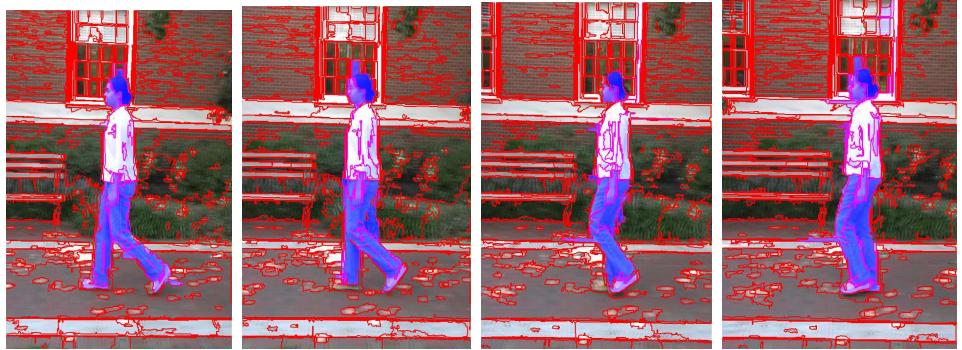


Figure 4.8: Another set of example frames for tracking with a moving camera. The outdoor scene contains more clustered foreground/background than *Karsten.avi*, and our segmentation results are less robust. To demonstrate the fast subject and camera motion in this sequence, note that these 4 frames last a quarter of second. The red pixels are on the boundary of segments; the tracked image segments associated with the foreground walking person is coded in blue. *The subject's white shirt is also correctly tracked. It does not appear in blue because its blue channel is already saturated.*

automatic learning. For surveillance video tracking, our results are very competitive with the state-of-art [?, 96] under even more challenging conditions.

Our approach does not depend on an image segmentation algorithm that totally respects the boundaries of the foreground object. Our novel bidirectional consistency check and resampling process has been demonstrated to be effectively robust and adaptive. We leave the explorations on supervised dimension reduction and density modeling techniques on image patch sets, optimal random patch sampling strategy, and self-tuned optimal image patch size searching as our future work.

In this paper, we extract foreground/background by classifying on individual image segments. It might improve the figure/ground segmentation accuracy by modeling their spatial pairwise relationships as well. This problem can be further solved using generative or discriminative random field (MRF/DRF) model or the boosting method on logistic classifiers [105]. In our current implementation, we treat online binary classification at each time step as an independent process, and plan to investigate on more efficient sequential learning methods. Finally, we focus on learning binary dynamic appearance models by assuming figure/ground are somewhat distribution-wise separable. Other cues, as object shape regularization and motion dynamics for tracking, can be combined to improve performance.

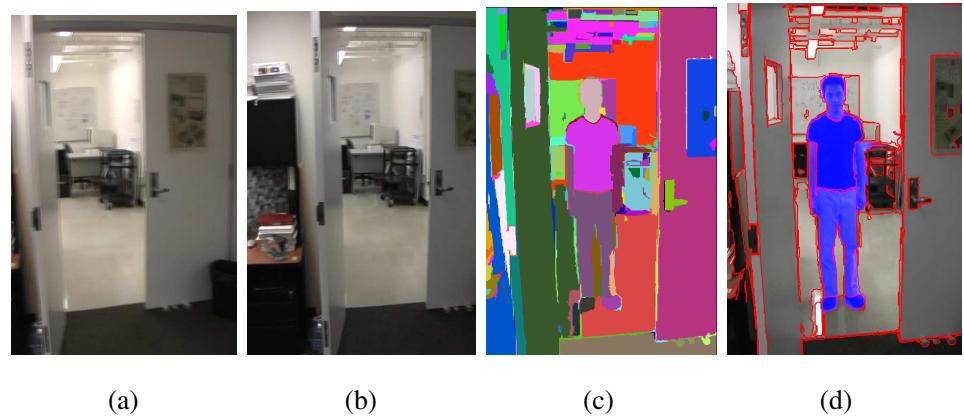


Figure 4.9: (a,b) 2 out of 12 background images; (c) the segmentation result for a testing image; (d) the testing image's detected foreground coded in blue.

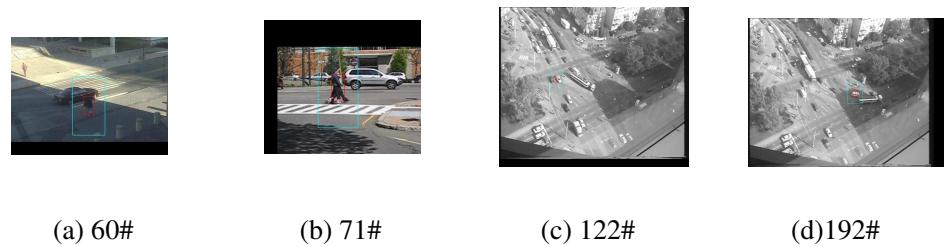


Figure 4.10: Sample tracked frames from four surveillance videos (a) Walking people tracking with low contrast (b) walking people tracking with handheld camera (c) car tracking in gray-level video; (d) car tracking with occlusion, self-turning and large illumination changes.

# Chapter 5

## Future Work

### 5.1 Future Work on Scene Recognition

#### 5.1.1 Linear and Nonlinear Discriminative Learning [84, 87, 69, 100, 168, 113, 132]

Learning a discriminative projection of data categories (such as image patches sampled from different material classes) is a general and important machine learning issue for classification problem. The discriminative projection reduces the data dimensionality and more importantly increases the separating margins between different data distributions. *Fisher/Linear Discriminant Analysis* (LDA) is the most popular algorithm which provides the optimal projection of discriminative feature extraction under the assumption that the distribution of data samples from each category is a Gaussian function with equal prior and covariance matrix [165]. This strict condition on the optimal projection assumption largely limits its application on real computer vision tasks, as many visual data distributions are believed to be highly non-linear and non-Gaussian.

As a non-linear extension of LDA, *Non-parametric discriminative analysis* (NDA) [87] is proposed and improved [69] for classification problems to overcome the Gaussian distribution constraint. The algorithm only involves computing covariances from local neighbors and the global distribution can have an arbitrary shape. Another important advantage of NDA is that the projected discriminative subspace is not limited to be equal or less than  $(c - 1)$  where  $c$  is the number of data classes. This property is very important to extract statistically sufficient features with large enough dimensions when  $c$  is too small. However NDA has an

extra requirement on finding the within-class and between-class nearest neighbor sets for each data, which can be very computationally expensive. To address this problem, *Locality-sensitive hashing* (LSH) [89, 107] provides a solution for efficient, approximate nearest neighbor searching with the bounded error tolerance and computational load. LSH has been successfully used for information retrieval over large collections of images and shapes [94, 93, 107]. The difficulty of employing LSH for NDA is that NDA requires finding both the within-class and between-class neighbor sets for each data entity while it is not very straightforward to integrate different data class labels into the hash function designing and coding in LSH. As shown in the second part of this proposal, we present a fast approximate method to compute the within-class and between-class nearest neighbor sets for data samples by using unsupervised data clustering. Hastie [100] describes another nonlinear global dimension reduction method by combining only local dimension information which has the similar computational complexity with NDA.

*Kernel Fisher Discriminant Analysis* (KFDA) [132] is another popular nonlinear discriminative learning technique . KFDA is designed to solve Fisher’s linear discriminant in the induced kernel feature space, which yields a nonlinear discriminant in input space. Many kernel functions (such as Gaussian, Polynomial, Spline and so on) can be used to map data into kernel space with possibly infinite dimensions. KFDA has been applied to many pattern classification problems, eg. digits recognition, face detection. The main difficulty of directly employing KFDA into our image patch discriminative learning is that the large number of training patch samples from all 20 material classes causes the computation of KFDA’s *pairwise kernel matrix* intractable. Some sparse approximation techniques [120, 119] on kernel method should be considered to address this problem in the future plan. Another possible drawback of KFDA is that kernel method originally intends to unravel the problem of (0-1) binary classifications and it can be cumbersome when the data class number  $c$  is too large.

Kosinov et al. presents a *distance-based discriminant analysis* (DDA) method using *iterative majorization* technique [113]. It gives promising results on both binary and multiple-class image categorization tasks. This method is also eligible being combined with kernel reproducing projection [143, 159] (Kernel DDA) for better performance on modelling more complex visual data. Zhu and Hastie [168] recently

propose a generalized non-parametric feature extraction method based on maximization of *likelihood ratio* (LR) which claims to overcome the Gaussian assumption of LDA. Yet, its numerical optimization process involves gradients on several hundred parameters in high dimension, which makes impractical for real world problems. The last noticeable technique on discriminative learning is called *discriminative log-linear model* [111] whose parameters are learned using *generalized iterative scaling* method [77]. This model have been tested on problems of artificial data classification [111] and image categorization [79], with a more appealing performance than LDA.

In the future work, we plan to evaluate and compare the performances of LDA, NDA, KFDA, DDA and KDDA for recognition purpose on our 20 class image patch database. Additionally, we are also interested to explore whether the linear coefficients computed from LDA can be further optimized in term of the class-separation criterion by using stochastic optimization methods, such as Monte Carlo Markov Chain [134].

### 5.1.2 (Hierarchical) Bayesian Learning [81]

Fei-fei and Perona propose a novel generative Bayesian hierarchical model to learn and recognize 13 natural scene categories using *Latent Dirichlet Allocation* [64]. This approach provides a principled method to learning relevant intermediate representations of visual scene automatically and without supervision. The most significant difference of this approach is that there is no need of requiring experts to annotate image regions for training purpose, compared with previous work (including ours [11]) [114, 101, 116, 105, 148, 127, 151]. *Latent Dirichlet Allocation* [64] is originally designed to represent and learn text-document models. With the adaption to image analysis [81], the algorithm provides a principled probabilistic framework for learning models of image features using codewords. It provides a learning hierarchy where feature-level codewords can further construct intermediate-level themes and the image categorization process is performed by integrating out all intermediate-level hidden variables in a Bayesian fashion. In general, parameter learning and inference for *Latent Dirichlet Allocation* model is not computationally tractable due to parameter coupling, but can be achieved via Monte Carlo Markov Chain [64], Laplace approximation or Variational approximation [81]. A very challenging photo database including 13 natural categories is

used to testify the proposed Bayesian hierarchical model, and a satisfactory categorization performance is reported [81]. Additionally, this model is also shown to "be able to group categories of images into a sensible hierarchy, similar to what humans would do" [81]. Our plan is to evaluate this model using our photo database [11] for recognition performance comparison. It will be very interesting to explore how much gain this model can achieve with much heavier computations than our algorithm.

### 5.1.3 Generative-Discriminative Random Field (DRF) for Material-Class Image Segmentation [118, 116, 117, 155, 156, 101]

Our scene recognition algorithm [11] has been demonstrated to be robust for noisy local image patch matchings (towards the learned appearance models of 20 material classes) and need no requirement on segmenting local image regions as well. However our approach is capable to generate the image patch-wise labels of 20 material classes, as shown in Figure 3.4. Note that it is a difficult task for explicitly segmenting image regions according to each material class, due to the inherent image appearance ambiguity. For example, an image patch of "blue sky" can be visually indistinguishable from a water patch [116, 101], unless we can leverage some contextual information into the classification process. The general technique on modelling image spatial contextual interactions is *Markov Random Field* (MRF) [88, 122] which basically performs as a discontinuity-preserving smoothing process to integrate local neighboring information for robust labelling. The *Simulated Annealing* and *Monte Carlo Markov Chain* based MRF parameter learning/inference methods [88] is very computational demanding and time-consuming. Recently (*Loopy*) *Belief Propagation* [137, 85] and (*Max-flow/mini-cut*) *Graph-cut* [112] has been demonstrated to be two standard fast MRF solvers in [152] with good results. In this proposal, we employ *loopy belief propagation* algorithm [85] to re-estimate the image patch's likelihood probability according to each material class density model for better material detection results (Figure 3.4). To make the smoothing process of local observation measurements adaptive to the image contents themselves, data observations can be directly fused into either the pairwise potential functions in belief propagation [11] or the pairwise interaction energy term in graph-cut [112]. In summary, all standard *Markov Random Field* model is considered as a probabilistic generative model.

Material-class image segmentation problem is indeed a conditional classification task (ie. finding hidden labels conditionally on image observations), while generative framework (eg. *Markov Random Field*) expends efforts on modelling the joint distribution of image observations and their semantic labels which can be very computationally expensive or even intractable. On the contrary, discriminative framework models the conditional posterior probability directly which can possibly result in a simpler solution. As noted in [116, 117], "a potential advantage of employing the discriminative approach is that the true underlying generative framework can be quite complex even though the class posterior is simple". Recently, *Discriminative Random Field* (DRF) is proposed by [116], for image patch labelling task, based on the idea of *Conditional Random Field* (CRF) [118]. DRF and CRF are discriminative models that formulate the conditional distribution over labels (as hidden variables) using discriminative classifiers. The most common classifier used in [118, 116, 117] is the logistic regression classifier [165, 5]. Both the association potential and interaction potential functions in DRF utilize the log-linear model over labels on site-wise or pairwise observations. Because the evaluation of the partition function in DRF is NP-hard problem, the model parameters are actually learned using various approximate techniques, eg. mean-field or pseudo-likelihood; and the inference process can be performed using *Iterated Conditional Modes* [63] (ICM), *loopy belief propagation* [85] (for site-wise *Maximum Posterior Marginal* solution) or *max-flow/min-cut* [112] (for *Maximum A Posterior* solution). For details, refer to [116, 117, 118, 122].

Some other researchers also propose variations of CRF type method for image labelling [101] and object recognition [156, 155]. Particularly, He et al. [101] adopt a (product of experts) neural network framework [102] to compute the product of the local neighboring distributions as a single distribution for conditional labelling. The neural network parameters are learned from a set of annotated images using contrastive divergence [102]. Instead of logistic regression or neural network, Torralba et al. [156, 155] describe the *Boosted Conditional Random Field* model by applying *real boosting* [86, 74] method to learn the graph structure and local evidence of a *conditional random field* [118] additively for contextual-level object detection.

In summary, DRF offers a much flexible way to model more complex interactions using a learned classification function in the field model, compared with MRF by merely taking a predefined prior over interac-

tions. DRF is trained from a set of supervised data so that it is more task-driven for good results. Another advantage of DRF over MRF is that the underlying interactions can be formulated in many forms: pixel to pixel, region to region, segment to segment, object part to object part [117]. It greatly enhances DRF's modelling capacity for diversely visual data. In this proposal, we plan to investigate a simplified version of CRF for the material-class image segmentation problem. It involves representing images as a set of regularly or randomly sampled image patches; learning discriminative model-association functions; learning discriminative pairwise interaction/potential functions and using loopy or tree structured belief propagation over looped or star-shape random graphs for integrated conditional inference. Note that the local random graph structures needs to be learned [156] or simply searched [65] before any parameter learning or inference.

## 5.2 Future Work on Dynamic Foreground/Background Segmentation

### 5.2.1 Modeling Spatial Interactions Among Image Segments

In above, we propose to extract foreground/background image or video regions by classifying over individual image segments produced by a general segmentor [82]. To use contextual constraints, the accuracy of figure/ground segmentation can be improved by modelling image segment's spatial interactions as well. This problem is generally addressed by *Markov random field* (MRF) model [88, 122, 85, 112], *Discriminative/Conditional Random Field* [116, 117, 118] or the *boosting-additive method on logistic classifiers* [105, 86, 74].

For pairs of neighboring image segments under finer scales (eg. images shown in Figure 4.6), the *between-similarity measurement* are usually defined on the low-level image observations, such as the matching scores of their associated color or gradient histograms. On the other hand, the logistic regression function or alternative probabilistic discriminative classifiers can be trained from the joint feature vectors of pairs of spatially adjacent segments under coarser scales [116, 117], using a supervised approach. The pairwise joint feature pattern is directly encoded by learning the parameters of classifier functions<sup>1</sup>. In the future, we plan

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<sup>1</sup>For larger scaled indoor and outdoor images, many neighboring image segments of foreground or background are visually dissimilar. Thus the interactions based on image similarity will model the labelling process of image segments very independently.

to investigate how to define the similarity measurement of a larger clique of image segments (ie. beyond pairwise interactions).

Classification on hypotheses of constellations of image segments for labelling stability is the key contextual learning/inference issue. Firstly, within *Markov Random Field* representation, segment-model appearance matching responses are stated as data-model *association energy terms* and segment's pairwise similarities as neighboring *interaction energy terms*<sup>2</sup>. The figure/ground segmentation problem is then formulated as a binary partitioning of a graph of nodes (ie. image segments), which can be solved using the *max-flow/mini-cut graph-cut* techniques [112, 67] for *Maximum A Posterior* solution or (*loopy*) *Belief Propagation* [137, 85] for *site-wise Maximum Posterior Marginal* solution. Secondly, *Discriminative/Conditional Random Field* [116, 117, 118] employs different classifiers to model the *association and interaction energy* terms. A collections of discriminative classifiers (which can be defined separately) and their spatial interactions largely increase the modelling capacity and flexibility of possible energy formulations. Classifiers are discriminatively trained from a set of annotated data, which can make task-oriented image recognition tasks with possibly more prevailing performance than general-purpose generative models (such as MRF with general prior setting). Thirdly, [105] has shown that hypotheses for constellations of image segments can be directly evaluated through using *additive-boosting* algorithms [86, 74] to measure the constellation's intrinsic content coherence under the same semantic meaning. A large collection of images have shown being successfully interpreted as regions of horizontal ground, up-frontal man-made structure and sky in [105].

### 5.2.2 Boundary-Preserving Image Segmentation

In this part of proposal, experiments are performed by using a general-purpose image segmentor [82]. By adjusting the segmentor's smoothing convolution kernel, scaling parameters and the minimal segmentation size, we expect that images are moderately over-segmented and all the object-level boundaries can be pre-

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In this case, logistic regression function can work on the joint feature space to learn the joint contextual pattern directly, regardless that the spatial image segment neighbors are visually similar or dissimilar.

<sup>2</sup>Especially, we borrow the energy term names from Graph-cut framework [112].

served for further processing (as described in [?]). The image segmentation method used in [82] is a graph based partitioning algorithm based on local image color-texture observations. Because it is designed for general purposes, object-level boundaries can not always be detected, especially for image sequences captured under severe appearance changing and illumination conditions. Parts of foreground and background regions can group into a single partition due to their local homogeneity under strong shadow and motion blur. To enhance the performance of boundary-preserving image segmentation, we propose to generate new segmentation boundaries by thresholding belief maps (ie. the matching distance/density maps in Figure 4.5) produced by the learned foreground/background appearance models [148, 117]. It can be considered as a specified image segmentor<sup>3</sup> by using the domain knowledge directly. It can help detect object-level boundaries based on the high-level multi-modal<sup>4</sup> appearance model. Similarly, foreground/background's probability spatial prior maps based on kernel density evaluation in video sequences can also be treated as belief maps on finding object-level boundaries by using the shape constraint.

### 5.2.3 Uncertainty Measurement and Random Graph Belief Propagation

Though image boundaries detection can be enhanced by employing both the general-purposed image segmentors [82, 147] and domain-oriented belief maps thresholding [148, 117], there is no theoretical guarantee that all desirable object-level boundaries can be preserved during above processes. In order to address this problem, we plan to measure the confidence or uncertainty of decision making over individual image segments (ie. classifying image segments into foreground or background classes). One simple possibility is that the uncertainty value can be obtained from the variance of matching scores from image patches within each segment towards the foreground/background appearance models. For instance, if all image patches have very similar matching responses given a specific appearance model, the decision making process is considered to be very confident. Otherwise, if there are a lot contradict matching scores, the aggregated classification process over the whole image segment should be much less confident.

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<sup>3</sup>It is equivalent to run a general-purpose image segmentor on real-valued belief maps.

<sup>4</sup>The multi-modal appearance model can possibly convert nonhomogeneous image regions into homogeneous belief maps, which make it more convenient for the specific task-driven segmentation.

Based on uncertainty measurements of image segments, we plan to develop a heuristic to find segment candidates with suspicious (possibly incorrect) foreground/background labels. For each of those candidates, we will not label it as a single unit, but rather label all individual image patches within the segment instead. We further plan to formulate the patch labelling process using a *Markov random field* or *conditional random field* representation by leveraging contextual constraints. A brief introduction on parameter learning and inference algorithms on MRF or DRF is formerly described in section 5.1. Because we usually need to label hundreds of patches inside suspicious image segments, the computational efficiency turns out to be the key issue. Due to this reason, we propose to assemble each image patch with its spatial neighbors (and temporal neighbors from contiguous frames in video) and simplify the graph topology of field model to be "star-shaped", which can be efficiently learned and inferred using *tree-structure belief propagation* [137, 65]. The potential functions of messages in *belief propagation* can be defined according to the principles of MRF or DRF (section 5.1) respectively.

In summary, images segments with high confidences will be labelled as foreground/background directly (using the simple aggregation process described in section 4.4), while image segments with low confidences will be marked as "suspicious candidates" for further patch-wise labelling. This is an adaptive classification strategy by first solving easier problems with less computations, and then harder problems containing more ambiguities using more computations (as shown in above).

#### **5.2.4 Parametric or Non-parametric Density base Appearance Model**

In section 4.5, we compare the classification performances of object-level image mapping for eight algorithms. Based on this comparison, we adopt *Kernel based density estimator* (KDE) [106] to construct the nonparametric appearance density functions for foreground/background segmentation in videos, because of KDE's representative flexibility and computational efficiency [106]. In future work, we can use clustering methods [5, 165] condensing the number of representative data samples in *Kernel density estimator* [106], to further improve the computational speed. Note that the model selection problem of clustering techniques is not very critical in our case, though there is no general good solution for that. Here we mainly concern

the trade-off between the density approximation accuracy of KDE towards the true density functions and the computation gain obtained by condensing the number of kernels in KDE. The clustering model complexity can be conservatively larger than the number of modes of the underlying multi-modal density functions. We plan to investigate this issue in future work.

Because we represent any given image as a chunk of sampled image patches [?], it is equivalent to represent a single huge dimensional data item by using a distribution of data samples with much lower dimensions. This representation can offer the modelling flexibility on controlling the rigidity on how much or different new data can be integrated into the model when considering the temporal model updating process from videos. For example, for the dynamic foreground appearance model, we expect that there is no significant model shifting from one frame to its successive frame in a video sequence. If some new "foreground-labelled" image patch samples from the next frame is visually very different<sup>5</sup> with respect to the current model, they have more chance to be outliers. If we assume that the background appearances can be freely changing consequently, there will be no "outlier-rejection" process for background. This heuristic can be formulated into the sample's surviving probability during resampling. More apparently different patch samples has lower probability to survive through resampling; and vice versa. Further more, this "*model integrating with rejection*" strategy should be designed to be adaptive with the changing behavior of image contents in videos. Fast changing videos allow more difference tolerance, and slow changing videos allow less difference tolerance in the temporal appearance model updating process<sup>6</sup>.

### **5.2.5 Automatic Key-Frame Selection for Interactive Foreground/Background Segmentation**

In video based foreground/background segmentation applications, we only label the first frame manually (or using an automatic initialization procedure described in section 4.5), build an initial appearance model for

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<sup>5</sup>The difference intensity can be measured as the density value of a new patch sample in frame  $t$  evaluated by the density functions up to frame  $t - 1$ . Higher density value means less different or more consistent, and vice versa.

<sup>6</sup>A similar term in machine learning literatures is called "*learning rate*" which is a tradeoff between biases towards the history model or new contents.

foreground and background, then propagate this model into successive frames. For fast content changing videos (for instance, Karsten.avi in section 4.5 needs to be relabelled when unseen background appears from behind the walking person. Unexpected, severely appearance-changing organs come out very commonly in surgical videos.), it is very normal that some formerly unseen image contents consequently appear as parts of dynamic background. The temporal adaption of the appearance model rooting from the first video frame may be insufficient to recover temporal fast image content changes. To address this problem, we propose to automatically extract multiple key-frames from a given video sequence based on techniques of pairwise image matching [92, 93, 94, 107, 89, 95, 141] and spectral/graph partitioning [135, 136]. Note that we define "*key-frames*" as image frames containing mutually distinct visual contents and (probably) representing modes in the whole video appearance model. Therefore we expect the video key-frames to be automatically extracted containing significantly more amounts of appearance contents than just the first frame. This can help the initial appearance model trained from multiple key-frames be more representative and propagate more effectively in the temporal domain.

Particularly, we purpose to employ the *pyramid match kernel* approach [92] (which is based on the ideas of multi-resolution histogram pyramid for recognition [95] and *positive-definite kernel* [145]) to efficiently and effectively compute the matching scores of any given pair of image frames by matching two distributions from sets of sampled image patches. The other two related image matching techniques are locality-sensitive hashing [107, 89] and Earth Mover's Distance [141, 93, 94]. After obtaining the matching scores for all image frame pairs in video, we can either form them into the *pairwise Affinity matrix* and perform the *spectral clustering* [135] to find modes of clusters as key-frames, or convert them as a *weighted graph model* and execute the *graph partitioning* algorithm [136] to select key-frames. For these automatically extracted key-frames, we plan to assign foreground/background labels onto their manually selected image regions by using a publicly available image annotation tool: "*Label Me*" [157], for the initial foreground/background appearance model construction.

# Chapter 6

## Appendices

### 6.1 Appendix 1: Grey Level Cooccurrence Matrices: GLCM

Let us denote GLCM a  $N \times N$  matrix  $P_{i,j}$  where  $N$  is the quantized level of pixel intensity and  $i, j = 0, 1, \dots, N - 1$ . The diagonal elements ( $i = j$ ) all represent pixel pairs with no grey level difference; while the off-diagonal cells ( $i \neq j$ ) represent pixel pairs with dissimilarity  $|i - j|$  increasing linearly away from the diagonal. Therefore we have  $dissimilarity = \sum_{i,j=1}^{N-1} (P(i,j) \times |i - j|)$ . Furthermore  $ASM = \sum_{i,j=1}^{N-1} P(i,j)^2$  measures the uniformity of the distribution of GLCM.  $\mu_i = \sum_{i,j=1}^{N-1} (P(i,j) \times i|)$  and  $\mu_j = \sum_{i,j=1}^{N-1} (P(i,j) \times j|)$  are the means of the reference pixels or neighbor pixels. Similarly,  $\sigma_i = \sqrt{\sum_{i,j=1}^{N-1} (P(i,j) \times (i - \mu_i)^2)}$  and  $\sigma_j = \sqrt{\sum_{i,j=1}^{N-1} (P(i,j) \times (j - \mu_j)^2)}$  are the respective standard deviations, and  $correlation = \sum_{i,j=1}^{N-1} (P(i,j) \times (i - \mu_i)(j - \mu_j)) / (\sigma_i \times \sigma_j)$ . If the above means and standard deviations are calculated from symmetrical GLCM,  $\mu = \mu_i = \mu_j$  and  $\sigma = \sigma_i = \sigma_j$ . Finally the output of 5 Haralick features are  $\{dissimilarity, ASM, \mu, \sigma, correlation\}$  for each GLCM<sup>1</sup>.

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<sup>1</sup>Note that we choose the pair of reference and neighbor pixels according to 4 directions (45 degree each) and 1 or 2 pixel offsets. Therefore we have 8 GLCMs for any image patch which results in a 40 component feature vector.

## 6.2 Appendix 2: Linear Discriminant Analysis: LDA

The following objective function

$$J(\omega) = \frac{\omega^T \mathbf{S}_B \omega}{\omega^T \mathbf{S}_W \omega} \quad (6.1)$$

is maximized by solving a generalized eigenvector equation

$$\mathbf{S}_B \omega = \lambda \mathbf{S}_W \omega \quad (6.2)$$

where

$$\mathbf{S}_W = \frac{1}{M} \sum_{i=1}^C \sum_{j=1}^M z_{ij} (X_j - m_i)(X_j - m_i)^T \quad (6.3)$$

$$\mathbf{S}_B = \sum_{i=1}^C \frac{M_i}{M} (m_i - m)(m_i - m)^T \quad (6.4)$$

Denote that  $\mathbf{S}_B$  and  $\mathbf{S}_W$  are respectively named the between-class or within-class scatter matrix,  $x_j$  is a feature vector,  $m_i$  is the mean of class  $i$  and  $m$  is the global mean of the data  $X$ ,  $i = 1 \dots C$  is a class number ( $C$  is the total number of classes) and the binary membership function

$$z_{ij} = \begin{cases} 1, & \text{if } x_j \in \text{class } i \\ 0, & \text{otherwise} \end{cases} \quad (6.5)$$

The LDA algorithm firstly perform the singular value decomposition (SVD) of  $\mathbf{S}_W$

$$\mathbf{S}_W = \mathbf{U} \Lambda \mathbf{U}^T \quad (6.6)$$

then transform  $\mathbf{S}_B$  into

$$\mathbf{S}'_B = \Lambda^{-\frac{1}{2}} \mathbf{U}^T \mathbf{S}_B \mathbf{U} \Lambda^{-\frac{1}{2}} \quad (6.7)$$

and compute the eigenvectors of

$$\mathbf{S}'_B \mathbf{V} = \mathbf{V} \hat{\Lambda} \quad (6.8)$$

where  $\hat{\Lambda}$  is the diagonal matrix of eigenvalues of  $\mathbf{S}'_B$ . The optimal feature vectors  $\mathbf{Z}$  are therefore

$$\mathbf{Z} = \mathbf{A}^T \mathbf{X} \quad (6.9)$$

through the projected transform  $\mathbf{A}^T = \mathbf{V}^T \Lambda^{-\frac{1}{2}} \mathbf{U}^T$ . For dimension reduction, only the subset of eigenvectors  $\mathbf{V}$  and  $\mathbf{U}$  with large eigenvalues are used in the transform. The dimension of the LDA projected subspace is at most  $C - 1$ .

### 6.3 Appendix 3: Discriminative-GMM Algorithm

Linear Discriminant Analysis (LDA) is proposed to maximize the following objective function

$$J(\omega) = \frac{\omega^T \mathcal{S}_B \omega}{\omega^T \mathcal{S}_W \omega} \quad (6.10)$$

by solving a generalized eigenvector equation

$$\mathcal{S}_B \omega = \lambda \mathcal{S}_W \omega \quad (6.11)$$

where

$$\mathcal{S}_W = \frac{1}{M} \sum_{i=1}^N \sum_{j=1}^M z_{ij} (x_j - m_i)(x_j - m_i)^T \quad (6.12)$$

$$\mathcal{S}_B = \sum_{i=1}^N \frac{M_i}{M} (m_i - m)(m_i - m)^T \quad (6.13)$$

Denote that  $\mathbf{S}_B$  and  $\mathbf{S}_W$  are respectively named the between-class or within-class scatter matrix,  $x_j$  is a feature vector,  $m_i$  is the mean of class  $i$  and  $m$  is the global mean of the data  $X$ ,  $i = 1 \dots C$  is a class number and the binary membership function

$$\mathcal{L}_{ij} = \begin{cases} 1, & \text{if } x_j \in \text{class } i \\ 0, & \text{otherwise} \end{cases} \quad (6.14)$$

The LDA algorithm firstly perform the singular value decomposition (SVD) of  $\mathcal{S}_W$

$$\mathcal{S}_W = \mathbf{U} \Lambda \mathbf{U}^T \quad (6.15)$$

then transform  $\mathcal{S}_B$  into

$$\mathcal{S}'_B = \Lambda^{-\frac{1}{2}} \mathbf{U}^T \mathcal{S}_B \mathbf{U} \Lambda^{-\frac{1}{2}} \quad (6.16)$$

and compute the eigenvectors of

$$\mathcal{S}'_B \mathbf{V} = \mathbf{V} \hat{\Lambda} \quad (6.17)$$

where  $\hat{\Lambda}$  is the diagonal matrix of eigenvalues of  $\mathcal{S}'_B$ . The optimal feature vectors  $\mathcal{Z}$  are therefore

$$\mathcal{Z} = \mathcal{A}^T \mathcal{X} \quad (6.18)$$

through the projected transform

$$\mathcal{A}^T = \mathbf{V}^T \Lambda^{-\frac{1}{2}} \mathbf{U}^T$$

For dimension reduction, only the subset of eigenvectors  $\mathbf{V}$  and  $\mathbf{U}$  with larger eigenvalues are used in the transform. The dimension of the LDA projected subspace is at most  $C - 1$ .

After the projected data  $\mathcal{Z}$  is obtained, GMM parameters  $\{\pi, \mu, \Sigma\}$  are initialized as follows

$$\mu_i = \sum_{j=1}^M \mathcal{L}_{ij} x_j, i = 1, 2, \dots, \mathcal{N} \quad (6.19)$$

and  $\pi_i = 1/\mathcal{N}$ ,  $\Sigma_i$  is a  $\mathcal{D}$  dimensional identical matrix where  $i = 1, 2, \dots, \mathcal{N}$ . From the initial estimates,  $\{\pi, \mu, \Sigma\}$  are then re-estimated iteratively in an Expectation-Maximization [40] fashion. In the E-step,

$$\mathcal{W}_{ij} = \pi_i P(z_j; \mu_i, \Sigma_i) \quad (6.20)$$

where  $\mathcal{W}_{ij}$  is the membership function of  $x_j$  with class  $i$  and is normalized by

$$\mathcal{W}_{ij} = \frac{\mathcal{W}_{ij}}{\sum_{i=1, \dots, \mathcal{N}} \mathcal{W}_{ij}} \quad (6.21)$$

to satisfy the constraint  $\sum_{i=1, \dots, \mathcal{N}} \mathcal{W}_{ij} = 1$ . In the M-step, parameters  $\{\pi, \mu, \Sigma\}$  are computed for each mixture density function as follows

$$\pi_i = \frac{1}{M} \sum_{j=1, \dots, M} \mathcal{W}_{ij} \quad (6.22)$$

$$\mu_i = \frac{\sum_{j=1, \dots, M} \mathcal{W}_{ij} z_j}{\sum_{j=1, \dots, M} \mathcal{W}_{ij}} \quad (6.23)$$

$$\Sigma_i = \frac{\sum_{j=1, \dots, M} \mathcal{W}_{ij} (z_j - \mu_i)(z_j - \mu_i)^T}{\sum_{j=1, \dots, M} \mathcal{W}_{ij}} \quad (6.24)$$

When the EM algorithm converges (normally after 10 iteration in our case), we can update the within-class and between-class scatter matrices  $\mathcal{S}_{\mathcal{W}}, \mathcal{S}_{\mathcal{B}}$  from the membership function  $\mathcal{W}$  estimated in GMM. The updated  $\mathcal{S}_{\mathcal{W}}, \mathcal{S}_{\mathcal{B}}$  improve the performance of LDA to find more discriminative projections.

$$\mathcal{S}_{\mathcal{W}} = \frac{1}{M} \sum_{i=1}^{\mathcal{N}} \sum_{j=1}^M \mathcal{W}_{ij} (x_j - m_i)(x_j - m_i)^T \quad (6.25)$$

$$\mathcal{S}_{\mathcal{B}} = \sum_{i=1}^{\mathcal{N}} \pi_i (m_i - m)(m_i - m)^T \quad (6.26)$$

where  $m_i$  is the mean of class  $i$ ,  $m$  is the mean of all the data.

$$m_i = \frac{\sum_{j=1,\dots,M} \mathcal{W}_{ij} x_j}{\sum_{j=1,\dots,M} \mathcal{W}_{ij}} \quad (6.27)$$

$$m = \frac{1}{M} \sum_{j=1,\dots,M} x_j \quad (6.28)$$

As a summary, our algorithm contains 2 loops of iterations. The outer loop is the LDA-GMM iterations through the update of  $\mathcal{S}_W, \mathcal{S}_B$  by  $\mathcal{W}$ ; the inner loop is the EM iterations of GMM. In section 3.5, the convergency of discriminative-GMM algorithm is validated experimentally in terms of the increasing log-likelihood of the fitness of the model and data

$$\mathcal{LL} = \sum_{j=1}^M \sum_{i=1}^N \log (\pi_i P(z_j; \mu_i, \Sigma_i)) \quad (6.29)$$

and the decreasing ratios of the incorrectly clustered data under the fixed model complexity.

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