

Unsupervised Image Segmentation based on Clustering and Watersheds

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Abstract

Uns

1. Introduction

Image segmentation and contour detection are closely related problems that rely on the same premise: Given an image, find a partition (Grouping) of this pixels according to the boundaries that compose it. From this premise, it is possible to treat a segmentation problem as a contour detection and viceversa, this approach allows to propose and define several procedures to solve and evaluate this problem.

Before the introduction of supervised learning techniques to solve this problem, all the approaches and proposed solutions were restricted only to the pixel information present on the image (Unsupervised segmentation) which served as input to different algorithms that provided a higher order representation of the image (Superpixels) or even partial segmentation images. Of these aforementioned algorithms, it is worth to mention clustering-based methods, such as those based on classic generic clustering methods, such as K-Means, Gaussian Mixture Models, Hierarchical clustering (To be explained during the following section) and Mean Shift [3]. Other important methods represent the image as a directed graph with

weights in which a minimal cut between regions delimited by a contour is desired, an example of an algorithm present on this category corresponds to the Normalized Cuts framework proposed by Shi and Malik [4]. Finally, other method families try to represent the segmentation problem from a morphological perspective, in which the image is seen as an intensity surface that describes valleys and intensity peaks, in which, boundaries can be seen as inflection peaks, this is the base of the Watershed transform method [2].

Although all the unsupervised methods yield reasonable results that group the pixels of an image into disjoint regions, these methods underperform the performance presented by the modern state-of-the-art supervised segmentation algorithms, which render the previously approach only of interest if a superpixel representation of an image is desired, also, some of the unsupervised models are suitable to semi-automatic applications, such as interactive segmentations.

2. Materials and Methods

About the Implementation

A total of four unsupervised segmentation routines were implemented and subject to direct comparison by using the labels and ground truth masks defined by the evaluation methodology proposed for this purpose, by comparing the contours obtained by thresholding the boundaries

present between regions after clustering an image with the reference contours present on the dataset. Three out of four methods are based on classical general-purpose clustering methods, namely, K-means, Gaussian Mixture Models and Hierarchical segmentation. Finally, the last method is based on hierarchical watersheds with extended h-minima imposition.

With respect to the image input representation to each of the segmentation methods, it is possible to notice that due the difference between the implementation and purpose of the clustering methods and the morphological segmentation based on watersheds, the input representation may vary, to accomplish this task, the image was

evaluated over several colorspace (RGB, HSV and La*b*), however, to cluster this representation, it is necessary to unroll each pixel coordinate triplet (1) as a column vector, as described on (2), in this case, the spatial coordinates maybe added to enforce spatial segmentation coherence and reduce the number of disjoint regions labeled to the same cluster without being in the same spatial area. Analogically, to process the image by employing the watershed transform, the grayscale gradient was calculated by using the Sobel derivatives, the flooding procedure was done over the image representation on any colorspace, without any spatial information. Finally, the contour detection and thresholding of the segmentation was done using the Canny edge detector.

$$\begin{bmatrix} (R_{1,1}, G_{1,1}, B_{1,1}) & (R_{1,2}, G_{1,2}, B_{1,2}) & \cdots & (R_{1,n}, G_{1,n}, B_{1,n}) \\ (R_{2,1}, G_{2,1}, B_{2,1}) & (R_{2,2}, G_{2,2}, B_{2,2}) & \cdots & (R_{2,n}, G_{2,n}, B_{2,n}) \\ \vdots & \cdots & \ddots & \vdots \\ (R_{m,1}, G_{m,1}, B_{m,1}) & (R_{m,2}, G_{m,2}, B_{m,2}) & \cdots & (R_{m,n}, G_{m,n}, B_{m,n}) \end{bmatrix} \quad (1)$$

$$\left[\begin{array}{cc|c} R_{1,1} & R_{1,2} & R_{m,n} \\ G_{1,1} & G_{1,2} & G_{m,n} \\ B_{1,1} & B_{1,2} & B_{m,n} \\ 1 & 1 & m \\ 1 & 2 & n \end{array} \right] \quad (2)$$

The implementation of the watershed procedure is based on the OpenCV¹ Python bindings, with the regional minima extraction routines adapted from the Gala library², the clustering methods are based on the Sckit-learn³ implementations. Also, the centroid and means initialization for K-Means and GMM was based on the KMC²⁴ algorithm [1], all the routines were implemented on Cython, to im-

prove execution times and the efficiency of the evaluation procedure.

About the models

K-Means

In the present experiment, K-Means was employed to segment an image by conforming clusters from pixel information, the clusters means were calculated by using the ℓ_2 distance metric (Euclidean), which gives rise to spherical and convex color clusters, which can be spatially sparse or localized, depending on the inclusion of spatial information. To

¹<http://opencv.org>

²<https://github.com/janelia-flyem/gala>

³<http://scikit-learn.org/stable/>

⁴<https://github.com/obachem/kmc2>

segment an input image, for each color triplet, the label of the closest mean is assigned, finally, the segmented image is reshaped into the image original dimensions.

Gaussian Mixture Model (GMM)

The GMM procedure pretends to explain the input image⁵ as the set of points spanned by a set of K gaussian distributions. To fit the set of distributions, it is essential to find the maximum likelihood parameters of each gaussian, namely, its mean μ_k and its covariance matrix Σ_k . To maximize those parameters, the Expectation Maximization (EM) algorithm is employed, this routine employs the Mahalanobis distance to model the data, which implies that the distribution may present skewness and doesn't have to be spheric or symmetrical around each axis. The maximization of the a priori probability of the model given the data allows to assign a membership probability to each of the K distributions to each value. In practice, the label of the distribution which have the greater probability is chosen.

3. Results

To evaluate unsupervised methods applied to image segmentation, it was necessary to formulate a common evaluation ground, in which every method can be compared directly with other methods, to address this issue, a subset of eight images were sampled from the Berkeley BSDS Segmentation database, each image

$$Acc = 1 - \frac{\|y - \hat{y}\|}{\|y + \hat{y}\|} \quad (3)$$

However, the color information alone is not enough to get a reasonable segmentation image because this method tends to group distant regions that present a similar color value on any space, this result may be relevant for other applications such as image compression but in the context of image segmentation this may degrade the performance of

the method, to account for this limitations, it is possible to include the spatial coordinates to each input vector, however, this causes the appearance of spherical localized regions, which implies that the final result of the segmentation is close to a Voronoi pattern.

4. Conclusions

References

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⁵It is assumed that its values are normally distributed