## A FACE TRACKER TRAJECTORIES CLUSTERING USING MUTUAL INFORMATION

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### **ABSTRACT**

In this paper we propose an algorithm for face tracker's trajectories clustering. Our approach is based on the mutual information of the images and more precisely its normalized version (NMI). We make use of 2 color channels from the HSV space (Hue and Saturation) in order to calculate a 4D joint histogram and therefore calculate the mutual information. In this paper we also develop an algorithm where we apply robust heuristics and make use of a tracker information in order to diminish dimensionality and augment accuracy of our results. It is a supervised clustering algorithm which is therefore used (fuzzy c-means) in order to gather same trajectories and same faces together.

## 1. INTRODUCTION

Face clustering is an important application for semantics extraction on video and can be used in a multitude of applications in video processing. It can contribute in many ways, like determining the primordial actors or the creation of databases' references or dialog detection and many others. Until now some interesting algorithms have been proposed in [1]-[3], but most of them are based in calibrated face images from news or face recognition databases like [1]. Our approach exploits the capabilities of joint entropy and mutual information in order to classify face tracker trajectories images exported from face tracker like the one proposed in [4]. In [5] we have proposed a face clustering approach using only results from a face detector. In this paper we propose a more elaborated method where the use of tracker's information as well as some heuristics are manifesting better results than our previous work.

Mutual information (MI) is a novel and useful tool in order to find similarities between information. More concretely, MI is defined as the information that is shared between two distributions. Until now, MI is much exploited in bioinformatics application and serves many purposes in that field from DNA sequences categorization [6] to classification of proteins. In image processing MI is used, in many reprise, in image registration for medical images and gives relatively good results. A good review for the use of mutual information in medical imaging can be found in [7].

In [5] we have proposed a face clustering approach using only results from a face detector and mutual information on the intensity channel of the image. In this paper we propose a more elab-

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orated method where the use of tracker's information as well as some heuristics which are manifesting better results than our previous work. We also make use of a more than 1 channel for the mutual information calculatation.we investigate as well the possibility of using this tool for classification of face images in a more realistic content such as movies, where difficulties arise from the fast variations of illumination, scale, pose etc.

The remainder of this paper is organized as follows: in Section 2 a mathematical presentation of the mutual information and its normalized version are presented, as well as the description of heuristics and tracker's results integration. In Section 3 clustering algorithm is presented. In Section 4 we demonstrate results for a real movie case. Finally in Section 5 further work and conclusions are discussed.

### 2. MUTUAL INFORMATION FOR FACE CLUSTERING

Mutual information is defined as the information shared between two distribution. Let X and Y be two distributions. We define the joint entropy as:

$$H(X,Y) = -\sum (p(x,y)\log(p(x,y))) \tag{1}$$

where p(x,y) the normalized (summed to one) probability density function of the common information of distribution X and Y. In the same way, we define the Shannon's entropy for X and Y as:

$$H(X) = -\sum (p(x))\log(p(x)) \tag{2}$$

Therefore we can define the mutual information as:

$$I(X;Y) = H(X) + H(Y) - H(X,Y)$$
(3)

where this give use the final equation of mutual information:

$$I(X;Y) = \sum_{x} \sum_{y} p(x,y) \log \frac{p(x,y)}{p(x)p(y)}$$
(4)

I(X;Y) is a quantity that measures the mutual dependence of two random variables. If we use a logarithm with base 2, then the measure is in bit. This quantity needs to be somehow normalized in order to create a uniform metric between different images and therefor be used as a similarity measure. For this reason, we use the normalized MI, which is defined as the quotient of the sum of two entropies with the joint entropy of those two distribution.

$$NMI(X;Y) = \frac{H(X) + H(Y)}{H(X,Y)} \tag{5}$$

Is is also useful to notice that:

$$NMI(X;Y) = \frac{H(X) + H(Y)}{H(X,Y)}$$

$$NMI(Y;X) = \frac{H(X) + H(Y)}{H(Y,X)}$$
(6)

$$NMI(Y;X) = \frac{H(X) + H(Y)}{H(Y,X)} \tag{7}$$

But as we know from (1):

$$H(X,Y) = H(Y,X) \tag{8}$$

So,

$$NMI(X;Y) = NMI(Y;X) \tag{9}$$

A very detailed explanation of how this normalizes the mutual information can be found in [8].

As mentioned before mutual information is proven to be a good similarity criterion for many cases. In our case we use this criterion in order to create a similarity matrix from the input images. We calculate the 2D histograms of the HS space of each face image acquired from the detection/tracking process and therefore from the two images we calculate the joint 4D histogram H where:

$$T(h_1, s_1, h_2, s_2) = |\{(k, l) \in N_1 \times N_2 / A(k, l) = (h_1, s_1) \ B(k, l) = (h_2, s_2)\} (10)$$

where  $|\cdot|$  denotes the cardinality of a set, A, B are face images and,  $N_1$  and  $N_2$  are the width and height respectively of both images.

By defining the joint histogram that way, we have to admit that in order to calculate it, images have to be of same size. This means that one has to resize one image to the other's dimensions. In our approach, and in order to equalize big scaling interpolation issues, we define a mean bounding box which is calculated from all bounding boxes that the face detector provides to us. This approach shows better results than if we scale every pair of images forward the bigger or the smaller of them. So every image is scaled towards this mean bounding box before the mutual information calculation.

Another issue is the fact of anisotropic scaling. Detector's results are bounding boxes where typically width and height are not equal. In order to scale forward a mean bounding box problems arise when the two dimensions are not equal. To override this, we calculate the bigger dimension of the bounding box and then we are taking the square box that equals this dimension centered to the original's bounding box center. Less literally, let  $B = \{x_1, y_1, x_2, y_2\}$ be a bounding box. We define the width as  $x_2 - x_1$  and the height as  $y_2 - y_1$ . From the two dimensions we take the bigger one and stretch the other at that size. The resulting bounding box for B for width bigger than height (resp. height bigger than width ), will be:

$$B_{new} = \{x_1, y_1 - k, x_2, y_2 + k\}$$

$$(resp. B_{new} = \{x_1 + k, y_1, x_2 - k, y_2\})$$

$$(11)$$

where k equals  $\frac{(x_2-x_1)-(y_2-y_1)}{2}$ .

We have noticed that problems arise from scaling issues that involves detectors inaccuracy. This means that if the face is not correctly detected and the face image contains a big amount of the background then scaling is mismatching the two face images and results are inaccurate. In order to override this bottleneck, another processing step is made, which is inspired from registration algorithms and aim in maximizing the accuracy of our results.

Once we have put the detector's outputs in the same scale we calculate the NMI for different frames of the target face image. We vary the bounding box's width and height from 80% to 120% of the

initial mean bounding box, with a step of 5%. The aforementioned values are calculated experimentally. In this way, we are trying to eliminate scaling problems due to detector's errors. Finally, we take the maximum of the calculated NMIs between the two images.

As mentioned before, the movies' context is dominated from several difficulties in order to extract content information. The way our approach is using the mutual information is undertaking those problems. By using the scale variance within the detectors results and the point-to-point approach of the joint entropy we have succeeded to provide good results in a very complicated task. In [9] the problem is tackled based on a preprocessing of the image. Our approach is trying to avoid the preprocess and goes deeper in the mutual information properties to that end.

#### 2.1. Mutual Information Vectors

Our algorithm consists of creating a vector of MIs for every image. The dimension of that vector is equal to the size of the face detection results' data set. For every face image in the results set we calculate the NMI between this image and any other, and therefore we create a vector v. All those vectors results in an  $M \times M$  matrix (where M the cardinal of the set of all detections from a video sequence) where every row i of that matrix will be the NMI of the i-th detection with all other images.

$$S(i, j) = NMI(FaceImage_i, FaceImage_j)$$
 (12)

It is obvious that the elements of the diagonal will have value one, which is the normalized mutual information of a face image with itself and also the matrix will be symmetric w.r.t the main diagonal. The diagonal property of the matrix is a forward effect of the MI symmetry shown in eq. (9). Those properties are very helpful because they drastically intervene in the time complexity of the algorithm. By using those properties the time complexity is minimized by a multiplicative factor of 0.5 and an additive factor of -M.

## 2.2. Heuristics and Tracking Information integration

We have tested two methods in order to use the tracking information within our framework so as to generate better results than our previous work in [5]. The first approach is to heuristically modify the similarity matrix in a way that the face images within the same trajectories will have a mutual information value of one and also faces appearing in the same frame will have a mutual information of 0. Less literally, let S be the aforementioned similarity matrix extracted as mentioned earlier. We create the new (robust) similarity matrix as follow:

$$S'(i,j) = \begin{cases} 1, & \text{if } (X_i, X_j) \text{ same tracking trajectory} \\ S(i,j), & \text{if non of the above is true} \end{cases}$$

where  $X_i$  and  $X_j$  are two face images. The other approach consists of creating a similarity matrix from the tracking results. In this approach we calculate a statistical measure from the face's belonging to the same trajectory and therefore we create an  $N \times N$  similarity matrix as follow:

$$ST(T_k, T_l) = f(B(k, l))$$
(14)

where f is the statistics function (in our case the min,max,mean and median are used), and B(k,l) is the submatrix of  $S^\prime$  created from the cross entries of the trajectory k and l as follow:

$$B(k,l) = \{ S'(i,j) | i \in k \text{ and } j \in l \}$$
 (15)

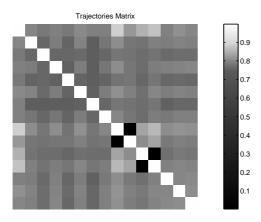


Fig. 1. Darker regions shows small amount of mutual information between tracker's results.

After we have created the matrix ST we apply a new heuristic rule in order to modify ST to a new matrix where we undertake the fact that two faces belonging to the same frame can not belong to the same person. This means that we have excluded cases where mirrors can create such an effect which are very rare. To do so we seek for the frames where two faces are located simultaneously and we put 0 at the cross reference cell of the ST matrix. In other words if a face belonging to trajectory 5 is in the same frame with a face belonging to trajectory 6 the the cell (5,6) of the matrix ST will be 0. The matrix ST after application of the heuristic rules is shown In figure 1. The reason we apply the heuristic rules is first in order to avoid faces belonging to the same trajectory to be clustered in to different groups and also to avoid faces that belong to the same frame to be clustered in the same group.

### 3. FUZZY C-MEANS CLUSTERING

Fuzzy c-means (FCM) is a method of clustering which allows one piece of data to belong to two or more clusters. This method initiated by Dunn in [10] and improved by Bezdek in [11] is frequently used in pattern recognition and it is based on minimization of an objective function. In order to cluster our results we use this algorithm which has been proven that in situation where we have a light mixture of classes' elements, it performs better than the simple k-means algorithm.

In order to use this algorithm we define every row of the similarity matrix S' as a different vector in an M-dimensional  $L^2$ -normed vector space over  $\mathbb{R}$ . In figure 2 and 3 one can see how those vectors are formed for two examples, one for 941-dimensional vectors with heuristics applied and another with 15-dimensional vectors from the tracker integration.

Therefore, we use the Euclidian distance to calculate distances between the vectors

$$dist(\mathbf{v_i}, \mathbf{v_j}) = \sqrt{\sum_{k=1}^{M} (v_{i_k} - v_{j_k})^2}$$
 (16)

and by those means to calculate a predefined number of clusters' centers. A detailed implementation of the FCM algorithm which was used, can be found in [12].

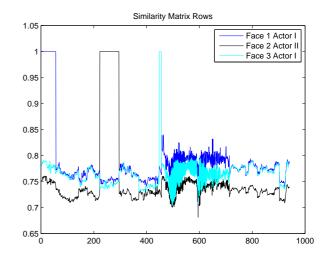


Fig. 2. 3 vectors which belong to 2 different actors from 3 distinct trajectories

Initialization has a significant role for FCM. So in order to provide better results the first centers are manually selected in a way that faces that corresponds in different actors constitutes a different initial center. A random selection of initial centers vary the results of a factor of 0.5% of false classification.

In the second approach the dimensionality of our vectors is drastically diminished due to the fact that M is now equal to the trajectories number and not to the face images cardinality. So from the 941 face images which results to a 941-dimensional space we arrive at a level of 15 dimension.

## 4. EXPERIMENT SCENARIO AND RESULTS

In order to test our algorithm we have conducted the following experiment. From a movie called "Two weeks notice" we have extracted a set of 941 detections which belong to 3 different actors, and also they are tracked 15 times. First we execute the detection algorithm in every first frame of a shot or if for some reason the tracker stops, then we redetect for that frame and retrack until a shot boundary is encountered.

The frames where selected so that light conditions and actor's poses vary, and also we have selected pieces of the film from different scenes. In the detector's results set we end up with a variety of face images in many poses and light conditions. With this approach we ensure the robustness of our algorithm in a multitude of different situations. In order to calculate the percentages of good and bad classifications we use a precision and recall like measure called F-measure [13]. The F-measure is calculated as follow: Let D represent a set and let  $\mathcal{C} = C_1, ..., C_k$  be a clustering of D. Moreover, let  $\mathcal{C}^* = C_1^*, ..., C_l^*$  design the human reference classification. Then the recall of cluster j with respect to class i, rec(i,j) is defined as  $\frac{|C_j \cap C_i^*|}{|C_i^*|}$ . The precision of cluster j with respect to cluster i, prec(i,j), is defined as  $\frac{|C_j \cap C_i^*|}{|C_j^*|}$ . The F-measure combines both values as follows:

$$F_{i,j} = \frac{2}{\frac{1}{prec(i,j)} + \frac{1}{rec(i,j)}} \tag{17}$$

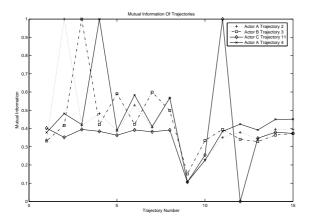


Fig. 3. 4 vectors which belong to 2 different actors from 4 distinct trajectories

Method	F-Measure
FCM on MI	65.4%
FCM on Robust MI	67.6%
FCM on Tracker MI using Min	66.33%
FCM on Tracker MI using Max	86.48%
FCM on Tracker MI using Mean	68.33%
FCM on Tracker MI using Median	60.0%

**Table 1**. Results Table of *F*-Measure.

The overall F-Measure of clustering is given by:

$$F = \sum_{i=1}^{l} \frac{|C_i^*|}{|D|} \cdot \max_{j=1,\dots,k} \{F_{i,j}\}$$
 (18)

We can easily note that a perfect fit between clustering and human reference leads to an F-measure score of 1. The F-measure is an external measure and thus it uses a human reference, i.e. it only shows how good the clustering is vis-a-vis to the human reference. The construction of the ground truth is mandatory for this process and unfortunately this kind of measure can not be used in real situations where human references are not available. Still, it is a very good measure for empirical evaluation of a new algorithm like the one proposed in this paper. In table 1 the results of the F-measure for all the experiments are illustrated.

# 5. CONCLUSIONS AND FUTURE WORK

We have developed a method for clustering face images within a very complex context such as movies. Results, as shown before, are rather promising for this difficult task if one considers the big variations that arise, w.r.t. light conditions, pose changes, emotions changes etc. in such a context. As face clustering has a lot of application in multimedia development, image processing and content based image retrieval applications (CBIR) we will investigate this problem further and we will concentrate our effort in the clustering process of the similarity matrix order to boost results.

The proposed method is a novel approach of the use that one can make of the mutual information in image analysis, and give good results in a hard task like the one we are solving. Exploration of the joint entropy and the mutual information on image data is shown to be a very good similarity criterion which can help in many other image processing application as well.

On the other hand, with our approach we minimize time complexity because of less preprocessing on the face images and the use of tracker information. This is an advantage for applications who needs fast clustering process, like interactive TV applications.

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