

GLOBAL JOURNAL OF COMPUTER SCIENCE AND TECHNOLOGY: F GRAPHICS & VISION Volume 14 Issue 3 Version 1.0 Year 2014 Type: Double Blind Peer Reviewed International Research Journal Publisher: Global Journals Inc. (USA) Online ISSN: 0975-4172 & Print ISSN: 0975-4350

## Image Segmentation for Animal Images using Finite Mixture of Pearson type VI Distribution

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GJCST-F Classification : 1.4.0



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# Image Segmentation for Animal Images using Finite Mixture of Pearson type VI Distribution

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Abstract- Image Segmentation is one of the significant tool for analyzing images, the feature vector of the images are different for different types of images. In remote sensing, Environmental ecological systems, forest studies. conservation of rare animals, the animal images are more important. In this paper we developed and analyze an image segmentation algorithm using mixture of Pearson Type VI Distribution. The Pearsonian Type VI Distribution will characterize the image regions of animal images. The appropriateness Pearsonian Type VI distribution for the pixel intensities of image region in animal images is carried by fitting Pearsonian Type VI Distribution to set of animal images taken from Berkeley image data set. The image segmentation algorithm is developed using EM algorithm for estimating the parameters of the model and maximum likelihood for image component under Bayesian framework. For fast convergence of EM algorithm the initial estimates of the model parameters are obtained by dividing the whole image into K image regions using K-means and Hierarchical clustering algorithm and utilizing the moment method of estimates. The performance of proposed algorithm is studied by conducting an experiment with set of animal images and computing image quality metrics such as PRI, GCE and VOI. A comparative study of developed image segmentation by Gaussian Mixture model and found the proposed algorithm performed better for animal images due to asymmetrically distributed nature of pixel intensities in the image regions.

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## I. INTRODUCTION

n image processing and retrievals image analysis plays a dominant role. The major task in image analysis is extracting useful information using features of the image. Generally image analysis techniques broadly grouping into groups namely (1) Structural methods (2) Statistical methods Raj Kumar et al (2011), among these two groups statistical methods are much popular. In Statistical methods one of the prime considerations is dividing whole image into different image regions using probability distributions. This type of method is usually referred as image Segmentation.

Much work has been reported in literature regarding image segmentation methods. Pal S.K and

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Pal N. R. (1993), Cheng et al (2001), Srinivasa et al (2007), Srinivas Y et al (2010), Prasad Reddy et al (2007) have reviewed the image segmentation methods. There is no unique image segmentation method available for analyzing all images. The image segmentation is basically dependent on type of images. The image broadly categorized into four types of categories . They are (1) Images on Earth (2) images of Humans and animals (3) images on sky (4) images on Water and (5) images of Nature. Among these categories the images of Human beings and Animals are in different in nature and features are associated with these images are different from others in some statistical sense. These images are Skewed in nature. Hence the image segmentation methods based on Gaussian mixture model given by Cheng et al (2001). Yamazaki T. et al (1998), Zhang Z.H et al (2003), Lie T. et al (1993) may not suit well. Even the methods given by Sesha sayee et al (2011), Srinivasa et al (2011) are also may not suit since these methods also focus on symmetry of the pixel intensities in the image region. Hence to have suitable and more appropriate image segmentation methods for animals, an image segmentation method using a mixture of Pearsonian Type VI Distribution is developed and analyzed. Here it is assumed that whole image is characterized by a mixture of Pearsonian Type VI probability model. The Pearsonian Type VI Distribution is skewed in nature having long upper tails. This distribution also includes several distributions as particular case. From the Berkeley image data set collected over animal images. It is evident that the pixel intensities of these images are well categorized by mixture of Pearsonian Type VI Distribution. The model parameters are estimated by updated equations of EM algorithm. The initial values of the model parameters of EM Algorithm are carried using Histograms of the whole image and K-means and Hierarchical clustering Algorithm and moment method of estimates. The image segmentation algorithm is developed through Maximum Likelihood component under Bayesian frame. The performance of image segmentation algorithm is skewed using image quality metrics and ground truth values. The comparative study of proposed algorithm with that of Gaussian Mixture Model is also carried.

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## II. MIXTURE OF PEARSON TYPE VI DISTRIBUTION

Usually the entire image is considered as a union of several image regions in low level image analysis and the image data is quantified by pixel intensities in each image region. Because of the fact that the brightness measured at a point in the image is influenced by various random factors like environmental conditions, vision, moisture, lighting etc, the pixel intensity z = f(x, y) for a given point (pixel) (x, y) is a random variable. It is generally assumed that the pixel intensities of the region follow a Pearson Type VI distribution in order to model the pixel intensities of the animal and human image regions. The probability density function of the pixel intensity is

$$f_{i}(z / a_{i1}, q_{i1}, q_{i2}) = \frac{\left(z_{s} - a_{i}\right)^{q_{i2}} \left(z_{s}\right)^{-q_{i1}}}{\left(a_{i}\right)^{(q_{i2} - q_{i1} + 1)} B(q_{i1} - q_{i2} - 1, q_{i2} + 1)} , (1)$$

$$a_{i} \le z_{i} < \alpha$$

The entire animal and human image is a collection of regions which are characterized by Pearson Type VI distribution. Here, it is assumed that the pixel intensities of the whole image follows a K – component mixture of Pearson Type VI distribution and its probability density function is of the form

$$p(z) = \sum_{i=1}^{K} \alpha_i f_i(z / a_{i1}, q_{i1}, q_{i2})$$
(2)

where, K is number of regions ,  $0 \le \alpha_i \le 1$  are

weights such that  $\sum \alpha_i = 1$  and  $f_i(z/a_{i1}, q_{i1}, q_{i2})$  is as given in equation (1). In the whole image  $\alpha_i$  is the weight associated with i<sup>th</sup> region. Usually the intensities of the pixel in the image regions are statistically correlated and can be reduced by spatial averaging ( Kelly P.A. et al (1998)) or spatial sampling (Lei T. and Sewehand W. (1992)). The pixels are considered to be uncorrelated and independent after reduction of correlation. The mean pixel intensity of the whole image is  $E(Z) = \sum_{i=1}^{K} \alpha u_i$ 

$$E(Z) = \sum_{i=1}^{n} \alpha_i \mu_i$$

## III. Estimation of the Model Parameters by EM Algorithm

In this section we derive the updated equations of the model parameters using Expectation Maximization (EM) algorithm. The likelihood function of the observations  $z_1, z_2, ..., z_N$  drawn from an image is

$$L(\theta) = \prod_{S=1}^{N} p(z_s, \theta^{(l)}) \text{ That is } L(\theta) = \prod_{S=1}^{N} \left( \sum_{i=1}^{K} \alpha_i f_i(z_s, \theta) \right)$$
  
This implies log  $L(\theta) = \sum_{S=1}^{N} \log \left( \sum_{i=1}^{K} \alpha_i f_i(z_s, \theta) \right)$ 

Where  $\theta = (a_{i1}, q_{i1}, q_{i2}, \alpha_i; i = 1, 2, ..., K)$  is the set of parameters.

$$\log L(\theta) = \sum_{s=1}^{N} \log \left[ \sum_{i=1}^{K} \frac{\alpha_i (z_s - a_i)^{q_{i2}} (z_s)^{-q_{i1}}}{(a_i)^{(q_{i2} - q_{i1} + 1)} B(q_{i1} - q_{i2} - 1, q_{i2} + 1)} \right] , (3)$$

The first step of the EM algorithm requires the estimation of the likelihood function of the sample observations.

a) E-Step

In the expectation (E) step, the expectation value of log  $L(\theta)$  with respect to the initial parameter vector  $\theta^{(0)}$  is

$$Q(\theta; \theta^{(0)}) = E_{\theta^{(0)}} \left[ \log L(\theta) / \overline{z} \right]$$

Given the initial parameters  $\theta^{(0)}$ , one can compute the density of pixel intensity  $z_{\rm x}$  as

$$p(z_s, \theta^{(l)}) = \sum_{i=1}^{K} \alpha_i^{(l)} f_i(z_s, \theta^{(l)})$$
$$L(\theta) = \prod_{s=1}^{N} p(z_s, \theta^{(l)})$$

This implies  $\log L(\theta) = \sum_{s=1}^{N} \log \left( \sum_{i=1}^{K} \alpha_i f_i(z_s, \theta^{(l)}) \right)$  (4)

The conditional probability of any observation  $z_{\rm s}$  , belongs to any region K is

$$t_{k}(z_{s}, \theta^{(l)}) = \left[\frac{\alpha_{k}^{(l)} f_{k}(z_{s}, \theta^{(l)})}{p(z_{s}, \theta^{(l)})}\right] = \left[\frac{\alpha_{k}^{(l)} f_{k}(z_{s}, \theta^{(l)})}{\frac{K}{\sum_{i=1}^{K} \alpha_{i}^{(l)} f_{i}(z_{s}, \theta^{(l)})}\right]$$

The expectation of the log likelihood function of the sample is

$$Q(\theta;\theta^{(l)}) = E_{\theta^{(l)}} \left[ \log L(\theta) / \overline{z} \right]$$

Following the heuristic arguments of Jeff A. Bilmes (1997) we have

$$Q(\theta;\theta^{(l)}) = \sum_{i=1}^{K} \sum_{s=1}^{N} \left( t_i(z_s,\theta^{(l)}) \left( \log f_i(z_s,\theta^{(l)}) + \log \alpha_i^{(l)} \right) \right)$$
(5)

But we have

$$f_i(z / a_{i1}, q_{i1}, q_{i2}) = \frac{\left(z_s - a_i\right)^{q_{i2}} \left(z_s\right)^{-q_{i1}}}{\left(a_i\right)^{(q_{i2} - q_{i1} + 1)} B(q_{i1} - q_{i2} - 1, q_{i2} + 1)}$$

$$Q\left(\theta;\theta^{(l)}\right) = \sum_{i=1}^{K} \sum_{s=1}^{N} \left( t_i(z_s,\theta^{(l)}) \left( \log f_i(z_s,\theta^{(l)}) + \log \alpha_i^{(l)} \right) \right)$$

b) M-Step

For obtaining the estimation of the model parameters one has to maximize  $\varrho(\theta; \theta^{(l)})$  such that  $\sum \alpha_i = 1$ . This can be solved by applying the standard solution method for constrained maximum by constructing the first order Lagrange type function,

$$S = \left[ E(\log L(\theta^{(l)})) + \lambda \left( 1 - \sum_{i=1}^{K} \alpha_i^{(l)} \right) \right]$$
(6)

where,  $\lambda$  is Lagrangian multiplier combining the constraint with the log likelihood function to be maximized.

Hence,  $\frac{\partial S}{\partial \alpha_i} = 0$ . This implies

$$\frac{\partial}{\partial \alpha_{i}} \left[ \sum_{i=1}^{K} \sum_{s=1}^{N} t_{i}(z_{s}, \theta^{(l)}) \left[ \log \left[ \frac{(z_{s} - a_{i})^{q_{i2}} (z_{s})^{-q_{i1}}}{(a_{i})^{(q_{i2} - q_{i1} + 1)} B(q_{i1} - q_{i2} - 1, q_{i2} + 1)} \right] + \log \alpha_{i} \right] \right] + \lambda \left( 1 - \sum_{i=1}^{K} \alpha_{i} \right) = 0$$

This implies  $\sum_{i=1}^{N} \frac{1}{\alpha_i} t_i(z_s, \theta^{(l)}) + \lambda = 0$ 

Summing both sides over all observations, we get  $\lambda = -N$  Therefore  $\hat{\alpha}_i = \frac{1}{N} \sum_{s=1}^{N} t_i(z_s, \theta^{(l)})$  The updated equation of  $\alpha_i$  for  $(l+1)^{\text{th}}$ iteration is  $\alpha_i^{(l+1)} = \frac{1}{N} \sum_{i=1}^{N} t_i(z_s, \theta^{(l)})$ 

$$\alpha_{i}^{(l+1)} = \frac{1}{N} \sum_{s=1}^{N} \left[ \frac{\alpha_{i}^{(l)} f_{i}(z_{s}, \theta^{(l)})}{\sum_{i=1}^{K} \alpha_{i}^{(l)} f_{i}(z_{s}, \theta^{(l)})} \right]$$
(7)

Therefore  $\frac{\partial}{\partial a_i} Q(\theta; \theta^{(l)}) = 0$  implies  $E\left[\frac{\partial \log L(\theta; \theta^{(l)})}{\partial a_{ij}}\right] = 0$ 

$$\frac{\partial}{\partial a_i} \left[ \sum_{i=1}^{K} \sum_{s=1}^{N} \left( t_i(z_s, \theta^{(l)}) \left( \log f_i(z_s, \theta^{(l)}) + \log \alpha_i^{(l)} \right) \right) \right] = 0$$

The updated equation of  $a_i$  at (l+1)th iteration is

$$a_{i}^{(l+1)} = \sum_{s=1}^{N} \left[ \frac{-(q_{i2}^{(l)} - q_{i1}^{(l)} + 1)(z_{s} - a_{i}^{(l)})t_{i}(z_{s}, \theta^{(l)})}{t_{i}(z_{s}, \theta^{(l)})q_{i2}^{(l)}} \right]$$

For updating the parameter  $\,q_{i1}\,$  , i = 1, 2, ..., K we consider the derivative of  $Q(\theta; \theta^{(l)})$  with respect to  $q_{i1}$  and equate it to zero. We have

$$Q(\theta; \theta^{(l)}) = E\left[\log L(\theta; \theta^{(l)})\right]$$
  
Therefore  $\frac{\partial}{\partial q_{i1}}Q(\theta; \theta^{(l)}) = 0$  implies  $E\left[\frac{\partial \log L(\theta; \theta^{(l)})}{\partial q_{i1}}\right] = 0$   
 $\frac{\partial}{\partial q_{i1}}\left[\sum_{i=1}^{K}\sum_{s=1}^{N}\left(t_i(z_s, \theta^{(l)})\left(\log f_i(z_s, \theta^{(l)}) + \log \alpha_i^{(l)}\right)\right)\right] = 0$   
 $q_{i1} = 1 - \sum_{s=1}^{N}\left[\frac{t_i(z_s, \theta^{(l)})}{\left[\left[\log \frac{a_i}{z_s} - \psi_0(q_{i1} - q_{i2} - 2 + 1) + \psi_0(q_{i1} - 1)\right]t_i(z_s, \theta^{(l)})\right]\right]$  (8)

The updated equation of  $q_{i1}$  at (l+1)th iteration is

$$q_{i}^{(l+1)} = 1 - \sum_{s=1}^{N} \left[ \frac{t_i(z_s, \theta^{(l)})}{\left[ \left[ \log \frac{a_i}{z_s} - \psi_0(q_{i_1}^{(l)} - q_{i_2}^{(l)} - 2 + 1) + \psi_0(q_{i_1}^{(l)} - 1) \right] t_i(z_s, \theta^{(l)}) \right]} \right] (9)$$

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For updating the parameter  $q_{i2}$ , i = 1, 2, ..., K we consider the derivative of  $Q( heta; heta^{(l)})$  with respect to  $q_{i2}$  and equate it to zero. We have

$$Q(\theta; \theta^{(l)}) = E\left[\log L(\theta; \theta^{(l)})\right]$$
  
Therefore  $\frac{\partial}{\partial q_{i2}} Q(\theta; \theta^{(l)}) = 0$  implies  $E\left[\frac{\partial \log L(\theta; \theta^{(l)})}{\partial q_{i2}}\right] = 0$   
 $\frac{\partial}{\partial q_{i2}} \left[\sum_{i=1}^{K} \sum_{s=1}^{N} \left(t_i(z_s, \theta^{(l)})\left(\log f_i(z_s, \theta^{(l)}) + \log \alpha_i^{(l)}\right)\right)\right] = 0$   
 $q_{i2} = \frac{\sum_{s=1}^{N} t_i(z_s, \theta^{(l)})}{\sum_{s=1}^{N} \left[\log \frac{(z_s - a_i)}{a_i} + \psi_0(q_{i1} - q_{i2} - 2 + 1) - \psi_0(q_{i2})\right]_{t_i}(z_s, \theta^{(l)})}$  (10)

The updated equation of  $q_{i2}$  at (l+1)th iteration is

$$q_{i2}^{(l+1)} = \frac{\sum_{s=1}^{N} t_i(z_s, \theta^{(l)})}{\sum_{s=1}^{N} \left[ \log \frac{(z_s - a_i)}{a_i} + \psi_0(q_{i1}^{(l)} - q_{i2}^{(l)} - 2 + 1) - \psi_0(q_{i2}^{(l)}) \right] t_i(z_s, \theta^{(l)})$$
(11)

Where 
$$t_i(z_s, \theta^{(l)}) = \frac{\alpha_i^{(l)} f_i(z_s, \theta^{(l)})}{\sum_{i=1}^{K} \alpha_i^{(l)} f_i(z_s, \theta^{(l)})}$$

## IV. INITIALIZATION OF THE PARAMETERS BY K - Means and Hierarchical Algorithm

Generally the efficiency of the EM algorithm depends upon the count of the regions in the image, during the estimation of the parameters. The number of mixture components taken for K – Means algorithm is, by plotting the histogram of the pixel intensities of the whole image. The number of peaks in the histogram can be taken as the initial value of the number of regions K.

The mixing parameters  $\alpha_i$  and the model parameters  $q_{i1}$ ,  $q_{i2}$  are usually considered as known apriori. Drawing a random sample from the entire image (Mclanchan G. and Peel D. (2000)) is the most commonly used method for initializing parameters. This method shows better performance, if the sample size is large and its computational time is heavily increased. When the sample size is small, there are some small regions which may not be sampled. To divide the whole image into various homogeneous regions we use the K – Means algorithm. In this algorithm the centroids of the clusters are recomputed as soon as the pixel joins a cluster.

#### a) K-Means Clustering Algorithm

It is one of the simplest clustering technique with a primary goal to find the partition of the data which minimizes the squared error or the sum of squared distances between all the points and their respective cluster centers (Rose H. Turi, (2001)). This K-means algorithm uses an iterative procedure and this procedure minimizes the sum of distances from each object to its cluster centroid, over all clusters. This procedure consists of the following steps.

- 1) Randomly choose K data points from the whole dataset as initial clusters. These data points represent initial cluster centroids.
- 2) Calculate Euclidean distance of each data point from each cluster centre and assign the data points to its nearest cluster centre.
- 3) Calculate new cluster centre so that squared error distance of each cluster should be minimum.
- 4) Repeat step 2 and 3 until clustering centers do not change.
- 5) Stop the process.

In the above algorithm, once only if all points have been allocated to their closed cluster centre then the cluster centers are updated. The advantage of this algorithm is that it is a very simple method, and based on intuition about the nature of a cluster, which is that the within cluster error should be as small as possible. The disadvantage of the K-Means algorithm is that the number of clusters must be supplied as a parameter, leading to the user having to decide what the best number of clusters for the image is (Rose H. Turi, (2001)). Success of K-means algorithm depends on the parameter K, number of clusters in image.

After determining the final values of K (number of regions) , we obtain the initial estimates of  $a_1a_2a_3a_4$  and  $\alpha_4$ 

 $a_i, q_{i1}, q_{i2}$  and  $\alpha_i$  for the *i*<sup>th</sup> region using the segmented region pixel intensities with Pearson Type VI distribution

. The initial estimate  $a_i$  is taken as  $a_i = 1/K$ , where i =

1,2,...,K. The parameters  $q_{i1}$  and  $q_{i2}$  are estimated by the method of moments as first moment  $\mu_1$  and its three central moments ( $\mu_2$ ,  $\mu_3$  and  $\mu_4$ ).

#### b) Hierarchical Clustering Algorithm

In order to utilize the EM algorithm we have to initialize the parameter  $\alpha_i$  and the model parameters  $q_{i1}$ ,  $q_{i2}$  which are generally considered as known apriori. The initial values of  $\alpha_i$  can be taken as  $\alpha_i = 1/K$  where, *K* is the number of image regions obtained from the Hierarchical clustering algorithm (Marr D. et al (1980)). The steps involved in hierarchical clustering algorithm are as follows.

Step 1: Start by assigning each item to a segment. Each of the N items, are associated with N segments, each containing just one item. Let the distances (similarities) between the segments be the same as the distances (similarities) between the items they contain.

Step 2: Find the closest (most similar) pair of segments and merge them into a single segment. The number of segments is now reduced by one. Compute distances (similarities) between the new segments and each of the old segments.

Step 3: Repeat steps 2 and 3 until all items are segmented.

Step 3 can be done in different ways, namely i) Single-Linkage ii) Complete-Linkage and iii) Average-Linkage segmenting. We consider the Average -Linkage methodology. Average-Linkage segmenting (also called the unweighted pair-group method using arithmetic averages), is one of the most widely used hierarchical clustering algorithms. The average linkage algorithm is obtained by defining the distance between two segments to be the average distance between a point in one segment and a point in the other segment. The algorithm is an agglomerative scheme that erases rows and columns in the proximity matrix as old segments are merged into new ones.

The proximity matrix is D = [d(i,j)]. The segments are assigned sequence numbers 0, 1..., (n-1) and L(k) is the level of the K<sup>th</sup> segment. A segment with a sequence number m is denoted by (m) and the proximity between segments (r) and (s) is denoted d [(r), (s)]. The algorithm is composed of the following steps: Begin with the disjoint segment having level L(0) = 0 and sequence number m = 0.

Find the average dissimilar pair of segments in the current segment, say pair [(r), (s)], for all pairs of segments in the current segment.

- 1. Increment the sequence number: m = m + 1. Merge segments (r) and (s) into a single segment to form the next segmenting m. Set the level of this segmenting to L(m) = d[(r), (s)].
- 2. Update the proximity matrix, D, by deleting the rows and columns corresponding to segments (r)

and (s) and adding a row and column corresponding to the newly formed segment. The proximity between the new segment, denoted (r, s) and old segment( K ) is defined in this way.

$$d_{(r,s) K} = \frac{\sum_{i} \sum_{j} d(i,j)}{N_{(r,s)} N_{K}}$$

where d(i, j) is the distance between object i in the cluster (r, s) and object j in the cluster K, and  $N_{(r,s)}$ and  $N_{(k)}$  are the number of items in the clusters (r, s) and K respectively. The above procedure is repeated till the distance between two clusters is less than the specified threshold value.

We obtain the initial estimates of  $q_{i1}$ ,  $q_{i2}$  and  $a_i$ for the i<sup>th</sup> region using the segmented region pixel intensities with the moment method given by Pearsonian Type VI distribution, only after determining the final values of K (number of regions). After getting these initial estimates, the final refined estimates of the parameters through EM algorithm given in section (III) is obtained.

## V. Segmentation Algorithm

In this section, the characteristics of the image segmentation algorithm are projected. After refining the parameters, the first step in image segmentation is allocating the pixels to the segments of the image. This operation is performed by Segmentation Algorithm which consists of four steps.

Step 1: Plot the histogram of the whole image.

Step 2: Obtain the initial estimates of the model parameters using K-Means algorithm and moment estimates for each image region as discussed in section IV.

Step 3: Obtain the refined estimates of the model parameters  $q_{i1}$ ,  $q_{i2}$  and  $\alpha_i$  for i=1, 2, ..., K using the EM algorithm with the updated equations given by (7), (9) and (11) respectively in section III.

Step 4: Assign each pixel into the corresponding  $j^{th}$  region (segment) according to the maximum likelihood of the  $j^{th}$  component  $L_{j.}$  That is

$$L_{j} = \max_{j \in k} \left[ \frac{\left( z_{s} - a_{i} \right)^{q_{j2}} \left( z_{s} \right)^{-q_{j1}}}{\left( a_{j} \right)^{(q_{j2} - q_{j1} + 1)} \beta(q_{j1} - q_{j2} - 1, q_{j2} + 1)} \right]$$
$$a_{i} \leq z_{i} < \alpha - \infty < q_{j1}, q_{j2} < \infty.$$

## VI. EXPERIMENTAL RESULTS

The performance of the developed a segmentation method for the natural images, which are considered on the earth. For implementing this algorithm, we need to initialize the model parameters, which are usually done by using moment method of estimations. Initially the feature vector is divided into

different segmented regions by making use of nonparametric methods of segmentation namely K-means algorithm and Hierarchical clustering algorithms.

An experiment is conducted with four images taken from Berkeley images dataset (http:// www. eecs. berkeley.edu/Research/Projects/CS/Vision/bsds/BSDS3 00/html). The images FACE, EAGLE, NEST BIRD and TIGER, are considered for image segmentation in order to demonstrate the utility of the image segmentation algorithm. The pixel intensities of the whole image are taken as feature. The pixel intensities of the image are always assumed to follow a mixture of Pearson Type VI distribution.

That is, the image contains K regions and pixel intensities in each image region follow a Pearson Type VI distribution with different parameters. The number of segments in each of the four images considered for experimentation are determined by the histogram of pixel intensities. The histograms of the pixel intensities of the four images are shown in Figure 1.



## Figure 1 : Histograms Of The Images

The initial estimates of the number of the regions K in each image are obtained and given in Table 1.

Table 1 : Initial Estimates Of K

IMAGE	TIGER	EAGLE	NEST BIRD	FACE
Estimate of K	2	3	4	4

From Table 1, we observe that the image TIGER has two segments, images EAGLE has three segments and images NEST BIRD AND FACE have four segments each. The initial values of the model parameters mi1,mi2, qi1,qi2 and  $\alpha$ i for i = 1, 2, ..., K, for each image region are computed by the method given in section III.

By making use of these initial estimates and the updated equations of the EM Algorithm given in Section III, the final estimates of the model parameters for each image are obtained and presented in Tables 2.a,2.b, 2.c, and 2.d for different images.

Estin	nated Values Numbe	Of The Paramete For of Image Regio	ers For TIGEI ons (K =2)	R Image		
Parameters	Estima Par	tion of Initial ameters	Estimation of Final Parameters by EM Algorithm			
	Imag	e Region	Imag	e Region		
	1-TYPEI	2-TYPEVI	1-TYPEI	2-TYPEVI		
$\alpha_i$	0.500	0.500	0.988622	0.011378		
$a_{i1}$	-97.4751	<i>a</i> <sub>i</sub> =453.1814	-0.00316	$a_i = 1$		
<i>a</i> <sub>i2</sub>	26.96539	$q_{i1} =$ 1.061354	-7.3732	$q_{i1} =$ 1.06892		
$m_{i1}$	0.783307	$q_{i2} = 0.061354$	-11.0395	$q_{i2}$ =0.062561		
$m_{i2}$	-0.21669	-	-23.8409	-		

#### Table-2.a

## Table-2.b

Estimated Values Of The Parameters For EAGLE Image Number of Image Regions (K = 3)						
Parameters	ers Estimation of Initial Parameters Estimation of Final Parameters by E					
		Image Region			Image Region	
	1-TYPEI	2-TYPEVI	3-TYPEI	1-TYPEI	2-TYPEVI	3-TYPEI
$\alpha_{_i}$	0.333	0.333	0.333	0.950792	0.00245	0.046758
$a_{i1}$	-42.6351	<i>a</i> <sub><i>i</i></sub> =-829.883	-35.4287	0.004741	a <sub>i =-4.01E-12</sub>	-1.2948
$a_{i2}$	38.46268	$q_{i1} = 1.027226$	38.6806	-0.1653	$q_{i1} = 1.028736$	14.15414
$m_{i1}$	0.525725	$q_{i2} = 0.027226$	0.47806	0.225525	$q_{i2} = 0.027134$	2.642036
$m_{i2}$	-0.47428	-	-0.52194	0.594823	-	0.405279

## Table-2.c

Estimated Values Of The Parameters For NEST BIRD Image Number of Image Regions ( $K = 4$ )								
		Estimation of Init	ial Paramete	ers	Estimation	n of Final Parame	eters by EM	Algorithm
Paramotora		Image Reg	gion			Image Re	gion	
Farameters	1-TYPEI	1-TYPEI 2-TYPEVI 3-TYPEI 4-TYPEI				2-TYPEVI	3-TYPEI	4-TYPEI
$lpha_{i}$	0.250	0.250	0.250	0.250	-0.03459	0.002342	0.650132	0.382119
$a_{i1}$	-24.3208	<i>a</i> <sub>i</sub> =181.3095	-15.9394	-30.2968	-1.38028	$a_i = 1$	-0.215	-0.26305
$a_{i2}$	28.31176	$q_{i1}$ =1.066072	11.42341	14.58499	12.98182	$q_{i1} =$ 1.073076	-0.19938	-0.68031
$m_{i1}$	0.462087 $q_{i2} = 0.066072$ 0.582521 0.675036 0.297129 $q_{i2} = 0.06951$ 2						2.420679	2.281604
$m_{i2}$	-0.53791	-	-0.41748	-0.32496	0.848453	-	0.469579	0.520931

Estimated Values Of The Parameters For FACE Image Number of Image Regions ( $K = 4$ )								
Parameters	Estimation of Initial Parameters Estimation of Final Parameters by EM						by EM	
		Imag	e Region			Image	Region	
	1-TYPEI	1-TYPEI 2-TYPEI 3-TYPEVI 4-TYPEI			1-TYPEI	2-TYPEI	3-TYPEVI	4-TYPEI
$\alpha_{_i}$	0.250	0.250	0.250	0.250	0.0068	0.9772	0.0078	0.0082
$a_{i1}$	-18.965	-24.176	<i>a</i> <sub>i</sub> = -468.346	-16.528	20.6072	-0.1116	<i>a</i> <sub>i</sub> =1.000	-2.290
$a_{i2}$	22.946	17.734	$q_{i1} =$ 1.047	18.557	493.753	-0.3840	$q_{i1}$ =-1.055	15.577
$m_{i1}$	0.4525	0.5768	$q_{i2} = 0.0471$	0.4710	-0.0283	2.4305	$q_{i2} = 0.0475$	2.664
$m_{i2}$	-0.5474	-0.423	-	-0.5289	-0.0819	0.4662	-	0.3995

#### Table-2.d

The probability density function of pixel intensities of each image is estimated by substituting the final estimates of the model parameters. The estimated probability density function of the pixel intensities of the image TIGER is



The estimated probability density function of the pixelintensities of the image EAGLE is



The estimated probability density function of the pixel intensities of the image NEST BIRD is



The estimated probability density function of the pixel intensities of the image FACE is



Using the estimated probability density function and image segmentation algorithm given in section III, the image segmentation is done for the five images under consideration. The original and

## segmented images are shown in Figure 2

Figure 2 : Original and Segmented Images



## VII. INITIALIZATION OF PARAMETERS BY HIERARCHICAL CLUSTERING ALGORITHM

In this section, we evaluate the efficiency of the proposed image segmentation algorithm. For this purpose of evaluation the images are collected from the Berkeley image data set. For this we need to randomly pick four images from the database and feature vector consisting of gray value for each pixel of the image. The feature vector in each image is modeled by using Pearsonian Type VI & Pearsonian Type I Distribution. By dividing all the pixel into different regions using Hierarchical Clustering Algorithm, the initial values of model parameters mi1, mi2, qi1,qi2 and  $\alpha$ i are obtained. Using these values and the updated equations of EM- Algorithm discussed in section III with MATLAB code, the final values of the model parameters are calculated and presented in the tables 3.a, 3.b, 3.c and 3.d

Estimated	Estimated Values Of The Parameters For TIGER Image Number of Image Regions (K =2)					
Parameters	Estimatio Paramet Hierarchica	n of Initial ters by I clustering	Estimatic Paramete Algo	on of Final ers by EM rithm		
	Image	Region	Image Region			
	1	2	1	2		
$\alpha_{i}$	0.500	0.500	2.01257	-1.01257		
$a_{i1}$	-75.717	-11.7563	-1.6508	-0.0459		
$a_{i2}$	82.970	740.670	88.3755	-28057.91		
$m_{i1}$	0.4771	0.0156	0.6101	40.2429		
$m_{i2}$	-0.5228	-0.9843	1.7071	0.01562		

#### Table-3.a

## Table-3.b

Estimated Values Of The Parameters For EAGLE Image Number of Image Regions (K =3)						
Parameters	Estimation of Initial Parameters by Hierarchical clustering Estimation of Final Parameters by EM					
	I	mage Regior	า		Image Regio	n
	1	2	3	1	2	3
$lpha_{i}$	0.333	0.333	0.333	0.4574	0.1921	0.3503
$a_{i1}$	-29.580	-99.0809	-41.2789	-0.1968	0.6095	-0.05994
$a_{i2}$	22.567	14.1523	35.7243	-0.7845	-0.70016	-1.0802
$m_{i1}$	0.5672	0.87501	0.53606	0.26831	2.08306	2.50880
$m_{i2}$	-0.4327	-0.1249	-0.4639	0.67742	0.61615	0.44170

#### Table-3.c

Estimated Values Of The Parameters For NEST BIRD Image Number of Image Regions (K =4)								
Estimation of Initial Parameters by       Estimation of Final Parameters by EM         Parameters       Hierarchical clustering						by EM		
Image Region					Image	Region		
	1	2	3	4	1	2	3	4
$lpha_{_i}$	0.250	0.250	0.250	0.25	0.8124	0.6205	-0.1252	-0.3077
$a_{i1}$	-29.742	-42.662	-22.721	-13.326	-0.39174	-0.2174	-0.5579	-0.0467
$a_{i2}$	22.299	18.108	27.072	75.188	-0.8320	-0.7572	17.451	1018.7
$m_{i1}$	0.5715 0.7020 0.4563 0.1506				0.31705	2.2480	2.7025	5.3865
$m_{i2}$	-0.4284	-0.2979	-0.5436	-0.8494	0.7971	0.5349	0.3905	0.14647

Estimated Values Of The Parameters For FACE Image Number of Image Regions ( $K = 4$ )								
Parameters	Estimation of Initial Parameters by Estimation of Final Parameters by EM Algorithm Parameters Hierarchical clustering							M Algorithm
	Image Region					Image	Region	
	1	2	3	4	1	2	3	4
$\alpha_{i}$	0.250	0.250	0.250	0.250	0.3736	0.2847	0.2526	0.0891
$a_{i1}$	-12.244	-12.556	-14.509	-68.801	-0.0611	-0.1155	-0.0811	1.5517
$a_{i2}$	71.505	11.0781	9.1209	57.659	2536.35	0.3296	0.1468	-5.4401
$m_{i1}$	0.1462 0.5312 0.6140 0.544				-0.3232	2.5188	2.3684	2.4925
$m_{i2}$	-0.8537	-0.4687	-0.3859	-0.4559	-2.3224	0.4387	0.4876	0.4465

#### Table-3.d

Substituting the final estimates of the model parameters, the probability density function of pixel intensities of each image is estimated.

The estimated probability density function of the pixel intensities of the image TIGER is



The estimated probability density function of the pixel intensities of the image EAGLE is



The estimated probability density function of the pixel intensities of the image NEST BIRD is



The estimated probability density function of the pixel intensities of the image BIRD is



The estimated probability density function of the pixel intensities of the image FACE is



Using the estimated probability density function and image segmentation algorithm given in section V, the image segmentation is done for the four images under consideration. The original and segmented images are shown in Figure 3.

#### Figure 3 : Original and Segmented Images



## VIII. Performance Evalution

In this paper we have conducted the experiment and also examined its performance by making use of the image segmentation algorithm. The performance evaluation of this segmentation technique is carried by obtaining the three performance measures namely, (i) probabilistic rand index (PRI), (ii) global consistence error (GCE) and (iii) variation of information (VOI). By computing the segmentation performance measures namely VOI, PRI and GCE for the five images under study using Pearsonian Type VI Distribution (PTVID-K), the performance of the developed algorithm is studied. The computed values of the performance measures for the developed algorithm and the earlier existing finite Gaussian mixture model(GMM) with K-means algorithm and Hierarchical algorithm are presented in Table 4 for a comparative study.

Table 4 : Segmentation Performace Measures

IMAGES	METHOD	PEF M	PERFORMACE MEASURES			
		PRI	GCE	VOI		
	GMM	0.8234	0.4956	2.568		
TIGER	PTVID-K	0.9896	0.4742	1.921		
	PTVID-H	0.9897	0.4762	1.920		
	GMM	0.8423	0.7006	8.354		
EAGLE	PTVID-K	0.8505	0.7109	7.577		
	PTVID-H	0.8627	0.7054	7.2002		

NECT	GMM	0.9793	0.9142	8.8837
BIRD	PTVID-K	0.0258	0.0124	6.7136
	PTVID-H	0.0074	0.0001	7.2132
	GMM	0.0201	0.0891	7.2546
FACE	PTVID-K	0.0223	0.0134	7.1556
	PTID-K	0.9559	0.8584	8.8772

From Table 4 it is identified that the PRI values of the existing algorithm based on finite Gaussian Mixture model for the five images considered for experimentation are less than the values from the segmentation algorithm based Pearsonian Type VI Distribution with K-means. Similarly GCE and VOI values of the proposed algorithm are less than that of finite Gaussian mixture model. This reveals the fact that the proposed algorithm outperforms the existing algorithm based on the finite Gaussian mixture model.

After developing the image segmentation method , it is required to verify the utility of segmentation in model building of the image for image retrieval. By subjective image quality testing or by objective image quality testing the performance evaluation of the retrieved image can be done. Since the numerical results of an objective measure allow a consistent comparison of different algorithms the objective image quality testing methods are often used. There are several image quality measures available for performance evaluation of the image segmentation method. An extensive survey of quality measures is given by Eskicioglu A.M. and Fisher P.S. (1995).

Using the estimated probability density functions of the images under consideration the retrieved images are obtained and are shown in Figure 4.

Figure 4 : The Original and Retrieved Images



For the above retrieved images FACE, NEST BIRD AND EAGLE The calculated image quality measures using proposed PTVID and GMM with Kmeans and Hierarchical algorithm are displayed in the Table 5.

Table 5 : Comparative S	Study of Image	Quality Metrics
-------------------------	----------------	-----------------

IMA	Ouality	GM	PTVI	PTVI	Standard
GE	Metrics	М	D-K	D- H	Limits
TIGER	Average Difference	0.483 7	0.420 3	0.419 3	Close to 0
	Maximum Distance	1.000 0	1.000	1.000 0	Close to 1
	Image Fidelity	1.000	1.000	0.999	Close to 1
	Mean Square	0 601	0.410	0.410	Close to 0
	Error	1	3	0.410	C103C 10 0
	Signal to	6.654	6.088	6.641	As big as
	Noise Ratio	2	5	6	possible
	Image Quality Index	$1.000 \\ 0$	1.002 5	1.002 5	Close to 1
EAGLE	Average	0.594	0.514	0.453	Close to 0
	Difference	6	1	8	
	Maximum	1.000	1.000	1.000	Close to 1
	Distance	0	0	0	
	Image	1.000	0.971	0.862	Close to 1
	Fidelity	0	7	8	
	Mean Square	0.494	0.348	0.236	Close to 0
	Error	6	7	1	
	Signal to	5.682	11.15	15.10	As big as
	Noise Ratio	8	1/	83	possible
	Image Quality	1.000	0.905	0.673	Close to 1
NEST BIRD	Average	0 493	0.409	0.377	Close to 0
	Difference	0.495	0.409	3	Close to 0
	Maximum	1 000	1 000	1 000	Close to 1
	Distance	0	0	0	close to 1
	Image	1.000	0.913	0.856	Close to 1
	Fidelity	0	6	846	
	Mean Square	0.493	0.445	0.378	Close to 0
	Error	0	3	3	
	Signal to	5.689	13.88	14.15	As big as
	Noise Ratio	7	03	35	possible
	Image Quality	1.001	0.731	0.571	Close to 1
	Average	1	0.317	0.221	Close to 0
FACE	Difference	0.579	0.517	7	
	Maximum	1.000	1.000	1.000	Close to 1
	Distance	0	0	0	
	Image	1.000	0.570	0.680	Close to 1
	Fidelity	0	3	4	
	Mean Square Error	0.507 9	0.455 6	0.165	Close to 0
	Signal to	5.625	19.67	20.75	As big as
	Noise Ratio	1	07	62	possible
	Image Quality	1.000	0.036	0.033	Close to 1
	Index	7	6	6	

It is perceived that all the image quality measures for the five images are meeting the standard criteria as given in the Table 5. Basing on the above mentioned quality metrics we can retrieve images accurately by using the proposed algorithm. A comparative study is done on the algorithm based on finite Gaussian mixture model with the proposed algorithm and it reveals that the MSE of the proposed model is less than that of the finite Gaussian mixture model. The performance of the proposed model in retrieving the images is better than the finite Gaussian mixture model by making use of these quality metrics.

## IX. Conclusion

In this paper, by using finite mixture of Pearsonian Type VI distribution a new model image segmentation is introduced and analyzed. The pixel intensities of animal images better characterizes the mixture of Pearsonian Type VI distribution which is validated through experiment with Berkeley image data set. The model parameters are estimated by using the EM Algorithm. By using the Maximum Likelihood estimates, the Segmentation Algorithm is developed under Bayesian framework. The Experiment on the Berkeley image data set reveals that this image segmentation method outperforms in segmenting the animal images then that of the existing algorithm basing on Gaussian mixture model with respect to image segmentation quality metrics such as PRI, GCE and VOI. The proposed algorithm is much useful for image analysis and image retrievals. The image that is developed can be extended with a K-dimensional feature vector for color images which will be takes as elsewhere.

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