3D model retrieval using meshSIFT descriptor and fuzzy C-means clustering

Najlaa Abd Hamza¹, Shatha Habeeb Jafer² and Raghad Mohammed Hadi³

¹College of Nursing, University of Baghdad, Iraq
²Computer Science Department, University of Technology, Iraq
³College of Administration and Economics, Mustansiriyah University, Iraq

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ABSTRACT

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Corresponding Author:

Najlaa Abd Hamza, College of Nursing, University of Baghdad, Bab Al-Muadham, Baghdad, Iraq. Email: najlaaabd@conursing.uobaghdad.edu.iq

1. INTRODUCTION

Retrieving desired models from databases in a short time has become a large problem for research because of the increase in the number of 3D models in company or internet databases [1]. The number of requests for high-quality three-dimensional models is increasing each day. To create an innovative, straightforward model, however, is a very time-consuming procedure. In cases where three-dimensional models can be retrieved professionally from a database, the work efficiency in associated fields is enhanced by reprocessing the model. There are two methods for the retrieval of three-dimensional models from a database: a technique based on key words, and a technique based on content [2, 3].

For content-based 3D model retrieval, features of a model should be found to describe it, called shape descriptors. Some of these descriptors are robust, but applying them in real time is time-consuming; others are less time-consuming but also less accurate [1]. In this paper, we propose using a meshSIFT descriptor [4] which is robust against missing data, large rotations and translations and can thus find a balance between speed and robustness.

In a content-based retrieval engine, an input model, termed query, is selected, and models which are similar to the query are retrieved from a model database. Applying the retrieval algorithm directly to 3D models in a database is time-consuming [2, 5]. In this paper we propose a pre-processing step as an attempt to reduce the time it takes to search for 3D models in a dataset, achieved by reducing the size of that dataset through undertaking a comparison between different 3D model features using a KLT algorithm. These main features are extracted by means of a meshSIFT descriptor, resulting in one 3D model from any similar group representing an

input to a Fuzzy C-Means clustering approach of partitioning the vertices of both dataset and query models. Final steps are then taken to calculate the similarity between the two 3D models (dataset and query).

Mohammadhassan Rezaei and Erkan Gunpinar [1] have proposed a method for 3D mesh retrieval which first divides a model into clusters by utilizing a geodesic distance metric, using a skeleton-based, K-means clustering method, and then computes its descriptor using the area of these clusters. Each cluster is signified by an area-based descriptor which is invariant to orientation and scale.

Xiaohong Liu et al. [2] have implemented view-based, three-dimensional methods of model retrieval. In their methods, bag-of-words (BOW), regularization-based SIFT features are obtained from three projected views of a three-dimensional model, before a distributed K-means clustering approach (based on Hadoop platform) is employed for the calculation of feature vectors and for the clustering of three-dimensional models. The model correspondence is computed using the distance between the query model and every cluster's center, and the cluster closest to the query model is returned as a model of retrieval.

Kuan-Sheng Zou et al. [6] have introduced a combined shape distribution (CSD) descriptor for the retrieval of three-dimensional models based on group integration and principal plane analysis.

Jing Zhang et al. [7] have proposed a new sketch-based 3D model retrieval procedure which employs skeleton characteristics as the features to define the object shape.

2. FUZZY C-MEANS CLUSTERING

Clustering can be defined as the technique of grouping a collection of data objects as clusters or classes in such a way that objects within a cluster are similar to one another, but dissimilar to the other clusters' objects [8, 9].

Fuzzy C-Means (FCM) is one of the highest public fuzzy combination methods. It was proposed by Dunn [10] in 1973 and then improved by Bezdek [11] in 1981. This algorithm is an iterative clustering technique which produces an optimal partition by reducing the objective function $J_m = (X, U, C)$ [12].

$$J_{m} = \sum_{i=0}^{n} \sum_{j=0}^{c} U_{ij^{m}} \left\| x_{i} - c_{j} \right\|^{2}$$

Where m denotes parameter of fuzziness which is used to modify the weighting effect of membership values, $U=[u_{ij}]$ is a matrix of fuzzy partition that denotes the belongingness degree (membership) for x_i into center c_i , this matrix must satisfy the following limitations, for each j, $1 \le j \le n: u_{ij} = 1$, and $||x_i - c_j||^2$ is the distance between sample x_i and center c_i [10].

The Fuzzy C-Means algorithm has been summarized in the stages below [8, 14]:

- a) Initialize matrix U= $[u_{ii}]$ with initial value $u^{(0)}$.
- b) For n-step: compute the matrix of the cluster center $Q^{(n)} = [c_i]$ with $u^{(n)}$.
- c) Update $u^{(n)}$, $u^{(n+1)}$.
- d) If $||u^{(n+1)} u^n|| < \varepsilon$ then finish, else return to step2.

where ε represents the termination state threshold.

2.1. MeshSIFT

The meshSIFT algorithm consists of three main stages [4]: detection of key-points, assignment of orientations and generation of the local feature descriptor. The following subsection gives a detailed account of each of these stages.

2.1.1. Key-point detection

This stage detects significant points on the mesh. As in SIFT [15, 16], a method of scale space is followed. In this scale space significant points are identified. Initially, a scale space is built holding smoothed input mesh versions, which are achieved with the approximation of a Gaussian filter for meshes. This approximation involves the following mesh convolutions with a binomial filter [4]:

$$X_{i} = \begin{cases} X \ i = 0\\ \widehat{M_{lt}} * X \ else \end{cases}$$

The scale space is constructed using the following formula [15]:

$$\widehat{DoM}_{l} = X_{i} - X_{i-1} = \left(\widehat{M}_{Jt} - \widehat{M}_{Jt-1}\right) * X$$

where X is the original mesh, \widehat{M}_{Jt} and \widehat{DoM}_i are respectively the approximations of Gauss and difference of Gauss. The Gaussian filter's standard deviation \widehat{M}_{Jt} , which has been approximated for finding a smoothed mesh X_i , has been given by Jt.

In order to discover significant points in the scale space, the mean curvature [15]

$$M(x,y) = \frac{(1+c_y^2)c_{xx}-2c_xc_yc_{xy}+(1+c_x^2)c_{yy}}{2(1+c_x^2+c_y^2)}$$

is subsequently calculated for every one of the vertices and at every one of the scales in the scale space, where M_i is a set holding the mean curvature for every one of the vertices at a scale i. Variances between consequent scales are calculated in the following way [4]:

$$DM_i = M_{i+1} - M_i$$

To identify the extrema of the scale space, the DM value at every one of the vertices is matched to the values of its neighbours, on the identical scale and on the lower and upper scales. A vertex is chosen only in that case where its value for DM is smaller or larger compared to all its neighbours. Lastly, the precise scale Jt is allocated to every scale-space extremum, directing to significant points (key-points) with an allocated scale [15].

2.1.2. Assignment of orientations

In this stage, every one of the key-points is allocated a recognized orientation, by stating each parameter in scale function Jt. The value of Jt around every one of the key-points is calculated. Initially, for every one of the vertices inside this area, the normal vector is calculated and the geodesic distance to the corresponding key-point is determined. Subsequently, all computed normal vectors are projected onto the tangent plane to the mesh X_i holding the key-point [4].

2.1.3. Local descriptor

The local descriptor now offers for every one of the key-points (with allocated scale and recognized orientation) a vector of the feature containing a sequence of concatenated histograms. Every one of those histograms has been intended over a minor circular area. In each region two histograms are computed where each one of them has 8 bins: one holding the index of the shape,

$$I(x,y) = \frac{2}{\pi} \tan^{-1} \left[\frac{s_1(x,y) + s_2(x,y)}{s_1(x,y) - s_2(x,y)} \right]$$

(where s₁ and s₂ are the maximum and minimum curvatures respectively), and the other holding the slant angles (that is, the angle of the projected normal and the recognized orientations between each one). Initially, every item for both histograms is Gauss-weighted with the geodesic distance to the key-point (J = 4.5. Jt) and Gaussian weighted with the geodesic distance to the region's centre (J = 4.5. Jt). Every histogram is then normalized and cropped to $\frac{1}{\sqrt{m}}$ (where m is the number of bins), decreasing the effect of big histogram values. In a last stage all histograms are concatenated [4]:

$$\widehat{VF_i} = (\widehat{his_S})(his_{slant\ angle}), i = 1, ..9$$

$$VF = (\widehat{VF_1}, \widehat{VF_2}, ..., \widehat{VF_9})...$$

where VF is the vector of final feature [4].

3. 3D Mesh Model Representation

A 3D mesh *m* signifies the 3D surface, employing groups of mesh components – vertices *E*, edges *G*, and polygons (facets) *T* – where m = (E, G, T). The mesh vertices signify 3D locations, $E \subset R^3$. Every edge, $g_i \in G$, is separated through two isolated vertices, $G \subset \{g_i = \{e_j, e_k\} \mid e, e_k \in E, j \neq k\}$. Where every facet, $t_i \in T$, is unique at the smallest three edges with the end area, each pair of edges share a vertex [17].

In the state of a triangular mesh (which is the most used kind of mesh because of its relative straight-forwardness), the facet t_i is exactly categorized by three edges.

 $T \subset \{t_i = \{d_m, d_b, d_p\} | d_m, d_b, d_p \in D, m \neq b,$ $m \neq p, \forall d_1, d_2 \in g_i \exists t (t \in d_1 \land t \in d_2)\}.$ Every facet is distinct through three different vertices, $T \subset \{t_i = \{e_m, e_b, e_p\} \mid e_m, e_b, e_p \in E, m \neq b, m \neq p, b \neq p\}$ [18]. The mesh is part of a model, and defines the object vertices. Figure 1 shows the key components of each mesh [19]. There are several kinds of mesh representations, which vary depending on how the mesh data are structured and stored: data as polygon mesh, vertex-polygon mesh, simple adjacency mesh, partial adjacency mesh, triangular mesh and winged-edge mesh [20].

The triangular mesh is the most generally-used mesh representation because of its simplicity. It usually stores the triangles and the vertices, with guides to all neighbouring triangles and vertices, in isolated tables. The information from the edge is understood and can be determined at any triangle by its edges or vertex (distinct from the adjacent vertices) [19].

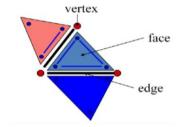


Figure 1. Example of vertices, faces, and edges

4. Kanade Lucas Tomasi (Klt) Algorithm

A KLT algorithm, which provides the greatest possible breadth, is used for feature tracking. KLT procedure has been presented by Lucas and Kanade, who suggest a process for registering two images for stereo correspondence. Their work has been extended by Tomasi and Kanade [21]. Firstly, the displacement of the tracked feature from one mesh to another is calculated. From this displacement calculation it is easy to compute the movement of the feature. KLT tracking procedure tracks the feature in two steps: it determines the traceable feature in the initial mesh before then tracking the identified features in the next mesh by utilizing the calculated movement [22].

Simple notification has been used to display the movement d of a feature from one mesh to the next. For ease the second mesh B(x) is redefined as B(x) = I(x) and the first mesh as A(x + d) = I(x + d). [22, 23]. The KLT tracker is the most efficient method of reducing

$$\epsilon(\mathbf{d}) = \frac{1}{2} \iint_{W} [J(\mathbf{x} + \mathbf{d}) - I(\mathbf{x})]^{2} \omega(\mathbf{x}) d\mathbf{x}$$

Compute the sum of squared error between mesh windows, then brute force search over d with normal cross correlation tracking. Here, I(x) is the value of the feature in the first mesh, J(x+d) is the value of the feature in the second mesh, x is the location of the feature in I, w(x) is a weight which is usually 1, and d is the movement of the feature.

The simple approach to tracking (also known as cross-correlation search) involves scanning a window around a search box (trying all values of d), and taking the one with the lowest cost, \in , above. This is surprisingly effective, but slow. If the search box has size 100×100 pixels, and each evaluation of \in involves a 15 x 15 pixel sum, then it takes at least $15 \times 15 \times 100 \times 100 = 2,250,000$ floating point operations per tracked feature per frame. The KLT takes a different approach. The idea is to reduce the remainder by differentiating $\in (d)$ with respect to d and taking a truncated Taylor expansion of J(x + d),

$$J(\mathbf{x} + \mathbf{d}) \approx J(\mathbf{x}) + \frac{\partial J(\mathbf{x})^T}{\partial \mathbf{x}} \mathbf{d}$$

where $\frac{\partial J(x)}{\partial x}$ is the image derivative in the x and y directions. Substituting this into the equation for \notin and setting derivatives to zero yields

$$\underbrace{\left[\iint_{W}\frac{\partial J(\mathbf{x})}{\partial \mathbf{x}}\frac{\partial J(\mathbf{x})^{T}}{\partial \mathbf{x}}\omega(\mathbf{x})d\mathbf{x}\right]d}_{Z\in R^{2\times2}}\underbrace{+\left[\iint_{W}[J(\mathbf{x})-I(\mathbf{x})]\frac{\partial J(\mathbf{x})^{T}}{\partial \mathbf{x}}\omega(\mathbf{x})d\mathbf{x}\right]}_{Z\in R^{2}}=Zd+e=0$$

Tracking thus reduces to repeatedly solving the simple 2×2 linear system Zd = e, where Z and e are readily computed from the images and gradients. The actual algorithm implemented is slightly more sophisticated, and uses multi-resolution image pyramids to enlarge the basin of attraction. By searching from the top of the pyramid down, it is possible to successfully track large motions [24].

5. System Overview

The proposed system uses three algorithms, as shown in Figure 2: Fuzzy C-Means clustering for 3D mesh models to partition the mesh into vertex clusters, a meshSIFT algorithm for deriving the key features of the 3D models, and a KLT algorithm for computing the similarity between the features in a pre-processing step. In the following sections, we give a brief overview of the process.

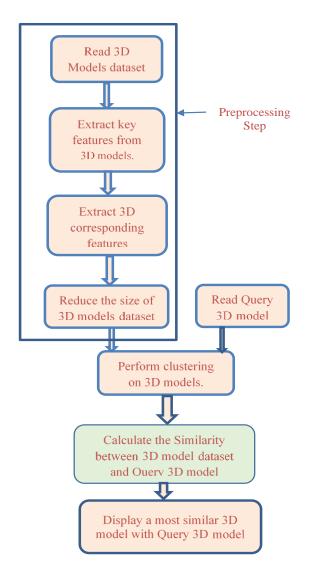


Figure 2. Block diagram of the proposed system

5.1. Pre-processing step

In this step, we attempt to reduce the size of the 3D model dataset by removing its redundancy. A detailed description of this step is provided in the following subsections.

5.1.1. Extracting the important features

In this stage, 3D model dataset is read and the key features are extracted for all 3D model datasets using the meshSIFT algorithm.

5.1.2. Extracting 3D corresponding features

In this step, the KLT algorithm is used to calculate the similarity between the features. After determining the features in the first 3D model, the algorithm tracks this feature with the other 3D models and then determines the most similar features between the 3D models. If the features between two models are equal (similarity 100%), then one model is rejected, and only one model from this group is taken.

The result of this step is a reduction in the size of the 3D models' dataset, decreasing the processing time by taking only one model from similar group models. Figure 3 shows our model database which contains 90 3D models from a Princeton shape benchmark [25] to authenticate the proposed 3D model retrieval method. When applying the pre-processing step to these 3D models, their number reduces from 80 to 60 by removing redundant models through taking only one from any similar group.



Figure 3. Model database

5.2. Reading the query 3D model

In this step, we read the query 3D model, a triangle mesh, before this model and all the 3D models in the dataset leftover after the pre-processing step enter clustering.

5.3. Clustering step

A Fuzzy C-means algorithm is now used to partition the mesh into vertex clusters. As previously outlined, this step is intended to create groups (i.e. clusters) of similar vertices according to similarity amongst the identical models. In the traditional FCM the centroid vector has been arbitrarily chosen for every one of the clusters, but in this paper, we propose to determine the centroid to get accurate clustering by taking the most featured vector (maximum value of feature) and determining it as centroid. In this work, we divided the model (dataset and query) into four clusters depending on shape descriptors. There are thus four clusters (similar in feature) to each model.

a)

5.4. Model retrieval

In this step, we calculate the similarity between the query 3D model and other 3D models. Similarity rate between two models are calculated based on the sum of squared differences (SSD) [6]:

 $Z = sum(m_1 - m_2)^2$

The individual steps involved in model retrieval are as follows:

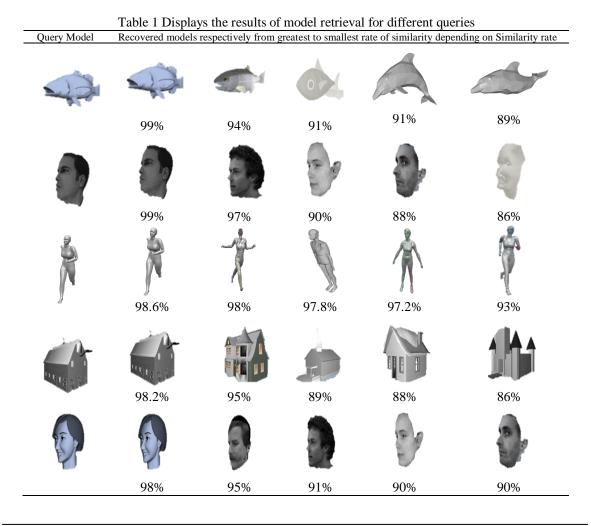
- Sort the clusters for each model from the smallest to largest.
- b) Compute the similarity between the largest cluster in the query 3D model and the largest cluster in the 3D model dataset.
- c) Sort the value of similarity from large to small.
- d) If the value is smaller than the threshold, reject the 3D model which contains this cluster.
- e) Calculate the similarity between the next largest cluster from the query 3D model and the other next largest cluster in the 3D model dataset, and go to step 3.

By applying these steps, we decrease the both the size of the 3D model data set and the time taken to process it.

6. RESULTS AND DISCUSSION

A set of testing models, comprising 80 3D mesh models from the Princeton shape benchmark [21], is used to certify the proposed algorithm. Retrieving is the procedure of discovering similar models to a query model. Table 1 shows the results of the proposed algorithm.

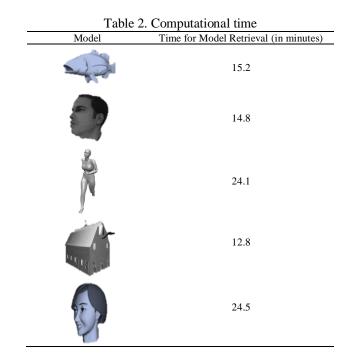
For a fish model, the most similar model is the fish model whose rate of similarity is 99%. For a face of women model, the most similar model is a women face model too. However, it has been observed that for a man face model, human model and a house model, the most similar models are the same model.



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According to these tests, it can be said that the proposed algorithm provides positive results in most cases. Computational time: time required for performing the proposed algorithm is shown in Table 2.



7. CONCLUSION

In the present paper, we propose an innovative technique for retrieving 3D models denoted by meshes. Having first extracted the key features of the models, the models are then separated into clusters by applying a Fuzzy C-means method depending on key features. Performance of the suggested retrieval method is shown via five experiments where rigid and non-rigid models are applied. According to the results, the suggested technique is successful in terms of its retrieving performance. Furthermore, it has been shown that the technique is invariant to different model poses, and therefore can be used for non-rigid models.

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BIOGRAPHIES OF AUTHORS



Najlaa Abd Hamza presently working as Assistant Professor of the College of Nursing, University of Baghdad, Iraq. She received Ph.D from Department of Computer Science, University of Technology, Iraq. Her Research interest includes Multimedia Processing, 3D Model processing, Feature Extraction, Security and Data Mining



Shatha Habeeb Jafer presently working as Assistant Professor & Head of the Information Systems Branch, Department of Computer Science, University of Technology, Iraq. She received Ph.D from Department of Computer Science, University of Technology, Iraq. Her Research interest includes internet of things, web of things, network management web application, Data Mining and security.



Raghad Mohammed Hadi presently working as Assistant Professor of the College of Administration and Economics, Mustansiriyah University, Iraq. She received Ph.D, from Department of Computer Science, University of Technology, Iraq. Her Research interest includes security, Network systems, AI, and Data Mining.