ABSTRACT

Realistic 3D modeling of human hand anatomy has a number of important applications, including real-time tracking, pose estimation, and human-computer interaction. However, the use of RGB-D sensors to accurately capture the full 3D shape of a hand is limited by self-occlusions, relatively smaller size of the hand and the requirement to capture multiple images. In this paper, we propose a method for generating a detailed, realistic hand model from a single frontal range scan and registered color image. In essence, our method converts this 2.5D data into a fully 3D model. The proposed approach extracts joint locations from the color image using a fingertip and interfinger region detector with a Naive Bayes probabilistic model. Direct correspondence between these joint locations in the range scan and a synthetic hand model are used to perform rigid registration, followed by a thin-plate-spline deformation that non-rigidly registers a synthetic model. This reconstructed model maintains similar geometric properties as the range scan, but also includes the back side of the hand. Experimental results demonstrate the promise of the method to produce detailed and realistic 3D hand geometry.

Index Terms— Hand model, pose estimation, reconstruction

1. INTRODUCTION

The recent introduction of real-time depth sensors and powerful computing devices has resulted in an increased research interest for natural interaction methods. The advent of mobile and wearable computers that utilize an ego-centric perspective have further necessitated the need for a novel interaction method that can overcome the limitations of the existing tangible and voice activated interfaces [1].

To this end, using the hand as an input for human computer interaction has undergone extensive research. A number of hand pose estimation methods have been proposed which require the use of synthetic hand models to reason how movements of a real hand result in corresponding variations in the input data [2, 3, 4, 5]. Furthermore the existing 3D reconstruction methods are limited by the noise, relatively smaller size of the hand and the requirement to capture multiple images for producing a complete hand model [6]. A realistic hand modeling method is required which can reconstruct a watertight realistic hand model encoding user-specific variations in hand geometry.

We present a method for generating detailed, realistic hand model from a single pair of frontal range scan and color image (as shown in Fig. 1). The proposed method extracts the location of joints using fingertips and valley points through a Naive Bayes probabilistic model. Direct correspondence between these joint locations on the range scan and the synthetic model are used to perform rigid registration, followed by a thin-plate-spline deformation that non-rigidly registers a synthetic model. This reconstructed model maintains similar geometric properties as the range scan, but also includes the back side of the hand. Experimental results demonstrate the promise of the method to produce detailed and realistic 3D hand geometry.

Previous work on 3D model construction using depth scanning has involved using a temporal sequence of depth and color images to reconstruct different surfaces [6]. In [7] a sequence of range images of the hand are used to build a user-specific hand model. Our method differs from these as we use a single front range and color scan of hand for modeling the geometry of the hand.

This paper is inspired by [8], which also produces 3D models from images. However, unlike [8] which only relies on color data, our method additionally considers a depth channel captured by a range scanner. The range data provides additional information regarding the shape of the hand, but only for the unoccluded frontal side. Our method detects creases using a probabilistic formulation and uses the range data as part of a 3D registration.
1.1. Contributions

To our knowledge, this is the first paper to produce a full watertight 3D hand model using only a frontal color and range image. In addition to the overall framework, we contribute a Naive Bayes approach to extract finger contours using gradient and spatial priors. The final result is a watertight synthetic hand model that captures the details in shape, size and pose of a real human hand. In addition, the proposed method implicitly infers the back side of the hand by applying the shape and size variations from the frontal scan.

2. METHOD

2.1. Problem Definition

Let \( S = \{I, D\} \) be a pair of frontal scans containing pairs of aligned color \((I)\) and depth \((D)\) images for different hands. We propose a method to estimate the 3D hand model \( M \) for each scan in \( S \). The estimated hand models vary in shape, size and pose, whereas the texture comes from an existing hand model and is common for all models in \( M \).

The overview of the proposed 3D hand model generation method is shown as a flowchart in Fig. 1. For each input image pair \( I \) and \( D \), the proposed method extracts the contour distance features \((CDF)\) using a segmented hand region [1]. Fingertips and valley points are then extracted from the \( CDF \) as local maxima and minima points. We then remove the excess wrist area from the hand by locating points on the radial and ulnar side of the palm and fitting them to an Euler spiral curve completion method [9]. A Naive Bayes probabilistic method is proposed to infer the finger joints' location which combines the probabilities of joint location from both the gradient image and an anatomical prior. A synthetic hand model with direct joint correspondence is then registered to a skeleton configuration acquired from finger joints in the real hand. Following this non-rigid registration is done using a thin-plate spline [10]. The final outcome is a complete 3D hand model with realistic shape, size and pose and a synthetic texture map on both the front and back. We discuss the proposed method in detail below.

2.2. Fingertips and Valley Extraction

\( CDF \) have been previously used for both pose and orientation estimation of the hand [1, 11]. We use these features as an initial step to localize the fingertips and valley points between the fingers. The fingertips \( F_i \) and valley points \( V_i \) are extracted from the \( CDF \) vector as the local maxima and minima, respectively, with each finger \( i \in \{\text{thumb, index, middle, ring, pinky}\} \). The proposed method uses an open hand pose, hence the number of visible fingertips and valley points remain constant.

In order to localize a finger completely we require the fingertip and two corresponding valley points on each side of the finger base. As only one valley point lies on the finger base for the thumb, index and pinky, we propose a method for extracting the additional base points on the radial and ulnar side of the hand, respectively.

Given the fingertip point \( F_i \), and the nearest valley point \( V_i \), a circle can be defined centered at \( F_i \) (shown in Fig. 2):

\[
(x - F_i^x)^2 + (y - F_i^y)^2 = r^2, \quad (1)
\]

where radius \( r \) is given as:

\[
r = \sqrt{(F_i^x - V_i^x)^2 + (F_i^y - V_i^y)^2}. \quad (2)
\]

Starting from a given valley point \( V_i \), and assuming the hand is roughly vertical in the 3D scan, we can grow the arc towards radial or ulnar side by decreasing or increasing the value of \( x \) in Eq. 2, respectively. The additional valley points \( V_j \) are extracted when the arc intersects with the hand contour, where \( j \in \{\text{radial-thumb, radial-index, ulnar-pinky}\} \).

2.3. Wrist Removal

In order for the proposed method to work across a range of different users, with varying wrist occlusions, we propose a method for wrist removal. This method uses an Euler spiral curve completion technique from [9, 11] and is similar to [8]. We extract two points on the ulnar and radial side of the hand using Eq. 2 for fingertip \( F_k \) where \( k \in \{\text{thumb, pinky}\} \). The radius \( r \) is increased by a scaling factor \( \alpha \), so that the arc intersects with the contour at points near the wrist. These points are then fed to the curve completion algorithm to remove the excess wrist area from both the hand depth \( D \) and color \( I \) image.

2.4. Joint Extraction

The analysis of the surface anatomy of the hand reveals its direct correspondence with the location of different skeletal
projection on medial axis. This figure is best viewed in color.

by gray horizontal lines, and (b) the corresponding gradient and the location of the anatomical prior from Eq. 4 depicted

puting the Euclidean distance between the fingertip

For a given finger

2.4.1. Crease detection

and the gradient of the frontal image of a given hand.

joints from (a) the real hand image, and (b) the rigidly regis-

Fig. 3: Crease detection using anatomical prior for the ring finger shows (a) fingertip

and synthetic hand model shows the location of fingertips and joints from (a) the real hand image, and (b) the rigidly regis-

tered synthetic model taken from [12].

joints. We utilize this observation to formulate the problem of joint localization by first locating the finger creases from the gradient of the color image. This is achieved using a Naïve Bayes probabilistic approach to combine the probability of the crease location from both the prior anatomical knowledge and the gradient of the frontal image of a given hand.

Fig. 4: Correspondence between joints extracted from the real and synthetic hand model shows the location of fingertips and joints from (a) the real hand image, and (b) the rigidly registered synthetic model taken from [12].

Finally the location of the crease is determined on the medial axis (Fig. 3(a) and (b)) as depicted by gray horizontal lines, and (b) the corresponding gradient projection on medial axis. This figure is best viewed in color.

Anatomical analysis of the hand reveals that the finger creases

2.4.1 Crease detection

For a given finger

and the location of the anatomical prior from Eq. 4 depicted

puting the Euclidean distance between the fingertip

For a given finger

we first estimate its length

\[ L_i \]

and the finger valley midpoint

\[ V_{mi} \]

in yellow, finger valley midpoint

\[ V_{mi} \]

in blue, the finger’s medial axis formed using

\[ F_i \]

and the location of the anatomical prior from Eq. 4 depicted

by gray horizontal lines, and (b) the corresponding gradient projection on medial axis. This figure is best viewed in color.

puting the Euclidean distance between the fingertip

\[ F_i \]

and the finger valley midpoint

\[ V_{mi} \]

in blue, the finger’s medial axis formed using

\[ F_i \]

and the location of the anatomical prior from Eq. 4 depicted

by gray horizontal lines, and (b) the corresponding gradient projection on medial axis. This figure is best viewed in color.

creases near

\[ h = 2 \]

and

\[ h = 3 \]

where

\[ \beta = 1/3 \]

relates the length

\[ L_i \]

of each finger

\[ i \]

anatomically to the most probable location of the creases.

The conditional probability

\[ p(h | c_i) \]

provides a region with high probability of finding a crease, however complete localization is achieved by formulating a gradient-based crease probability. The gradient of the color image is projected to each finger’s medial axis defined along the fingertip

\[ F_i \]

and the finger valley midpoint

\[ V_{mi} \].

This projection is achieved by accumulating the gradient output along the normal of medial axis for all points between

\[ F_i \]

and

\[ V_{mi} \].

The extracted signal is noisy due to the highly textured hand surface. We suppress this noise using a Gaussian filter with a standard deviation

\[ \sigma_f = 1.5 \].

This filtered signal represents the probability of crease location

\[ p(c_i) \]

in gradient of color image.

Assuming that both probabilities are independent, a Naïve Bayes method can be applied to achieve a combined probability

\[ p(c_i | h) \]

given by:

\[ p(c_i | h) \propto p(h | c_i) p(c_i) \cdot \] (5)

for each crease

\[ h \in \{1, 2, 3\} \]

as shown in Fig. 3.

Finally the location of the crease is determined on the medial axis by the maximum-a-posteriori (MAP) estimation for each crease in

\[ h \in \{1, 2, 3\} \].

\[ c_i* = \arg \max_{c_i} p(c_i | h) \cdot \] (6)

The extracted crease locations are then used to acquire the joint location from depth image. The interphalangeal joints (IP) directly lie under finger creases near

\[ h = 2 \]

and

\[ h = 3 \],

whereas the metacarpo-phalangeal joint (MP) lies 20% of finger length

\[ L_i \]

below

\[ h = 1 \]

on medial axis (Fig. 3(a) and Fig. 4(a)).

2.5. Rigid Registration

The proposed method utilizes direct joint correspondences of a synthetic hand model with the extracted joints of a real hand for rigid registration. The joint configurations from real hands are extracted in the form of joint angles, which are directly applied to the hand model. Fig. 4 shows the corresponding joints in both the real and synthetic hand model.

2.6. Non-Rigid Registration

The size and shape of the rigidly registered synthetic hand model differs from the real hand (Fig. 4). This requires non-rigid registration which is performed by building correspondences based on the location of the fingertips and finger valley points on the contour of both the range data and the
rigidly registered synthetic model. A thin-plate-spline deformation model is first built using these corresponding contour points which model the non-rigid transformation [10]. This is then applied to the synthetic hand model resulting in a reconstructed model of the real hand.

3. EXPERIMENTAL EVALUATION

We evaluate our method on the dataset from [13], which contains high resolution frontal 3D hand range and color scans acquired from different individuals. The error for each vertex on the generated 3D model is calculated by computing the nearest Euclidean distance, measured in mm, with the frontal range scan of the hand. Table 1 presents minimum, maximum and average error for 7 reconstructed models, whereas Fig. 6 localizes these errors for one of the model surfaces. The total average error of 3.65mm in Table 1 depicts the method’s ability to reconstruct the majority of the vertices with high accuracy. It can be observed that the majority of the errors exist in the inner palm and wrist region. This is due to the inability of our method to capture the complex shape of the palm and the exclusion of the wrist region from the modeling data. Another source of error is the thumb region, which arises due to the inability of our synthetic hand model to register the complex joint configurations of the thumb in real hands. Despite these limitations, overall the method is effective in reconstructing an accurate 3D hand model that matches the range data (Fig. 5).

In a separate experiment, the ability of the proposed method to automatically infer the back side of the hand is evaluated using a full 3D model of a person’s hand, providing a ground truth (GT) model. We use computer graphics to render this GT model and acquire frontal range and color images. These are then used in our proposed method to generate a 3D hand model. We then compare this reconstructed model to the original GT which is presented in Table 1 and depicted on model vertices in Fig. 7. Average error for both the front and the back is 2.51mm which is comparable to the average error for front only validation. Fig. 7 visualizes the error on the GT model, where larger errors follow our earlier observation for front side, particularly for the thumb, while the generated surface has no big error on the back side.

4. CONCLUSION

We proposed a method for generating realistic 3D hand models using only single frontal depth and color images of hands. This method used the fingertip and valley points and a probabilistic Naive Bayes approach to locate joints from the hand range scan and color image. The direct correspondences between the extracted joint locations with a synthetic model were used to perform rigid registration. This was followed by a thin-plate-spline deformation to produce a reconstructed model. The experimental evaluation showed an average error of 3.65mm. A GT hand model was also used to validate the accuracy of the inferred back side of the hand where average error for the whole reconstructed model was 2.51mm.

Table 1: Euclidean distance error between input data and Predicted model. All measurements are in millimeters (mm).

<table>
<thead>
<tr>
<th>Input data</th>
<th>Min error</th>
<th>Max error</th>
<th>Average error</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>0.04</td>
<td>25.02</td>
<td>5.58</td>
</tr>
<tr>
<td>2</td>
<td>0.10</td>
<td>10.54</td>
<td>2.74</td>
</tr>
<tr>
<td>3</td>
<td>0.03</td>
<td>20.47</td>
<td>2.65</td>
</tr>
<tr>
<td>4</td>
<td>0.07</td>
<td>12.20</td>
<td>3.43</td>
</tr>
<tr>
<td>5</td>
<td>0.03</td>
<td>16.06</td>
<td>3.69</td>
</tr>
<tr>
<td>6</td>
<td>0.08</td>
<td>15.06</td>
<td>2.58</td>
</tr>
<tr>
<td>7</td>
<td>0.09</td>
<td>19.29</td>
<td>4.85</td>
</tr>
<tr>
<td>Total Average</td>
<td>0.06</td>
<td>16.95</td>
<td>3.65</td>
</tr>
<tr>
<td>GT Model (Fig. 7)</td>
<td>0.12</td>
<td>13.91</td>
<td>2.51</td>
</tr>
</tbody>
</table>

Fig. 6: Error (a) in the hand model generation visualized using Euclidean distance between the input range and the generated model for model 5 in Table 1 and (b) the generated model.

Fig. 7: Euclidean distance error using a GT synthetic model with both the front and back side.
5. REFERENCES


