

3D HUMAN BODY POSE ESTIMATION BY SUPERQUADRICS

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Abstract: This paper presents a method for 3D Human Body pose estimation. 3D real data of the searched object is acquired by a multi-camera system and segmented by a special preprocessing algorithm based on clothing analysis. The human body model is built by nine SuperQuadrics (SQ) with a-priori known anthropometric scaling and shape parameters. The pose is estimated hierarchically by RANSAC-object search with a least square fitting 3D point cloud to SQ models: at first the body, and then the limbs. The solution is verified by evaluating the matching score, i.e. the number of inliers corresponding to a-priori chosen distance threshold, and comparing this score with admissible inlier threshold for the body and limbs. This method can be used for 3D object recognition, localization and pose estimation of Human Body.

1 INTRODUCTION

3D human body recognition and pose recovery are the important problems in computer vision and robotics with many potential applications including motion capture, human-computer interaction, sport and medical analysis, video surveillance, etc. The human body pose estimation from 3D real data obtained by a multi-camera system can be solved different ways. A generic humanoid model approximating a subject's shape can use either simple shape primitives (cylinders, cones, ellipsoids, and superquadrics) or a surface (polygonal mesh, sub-division surface) articulated using the kinematic skeleton (Forsyth, et al., 2005; Moeslund, et al, 2006; Balan, et al. 2007; Mun Wai Lee and Cohen, 2004; Ivecovic and Trucco, 2006). We consider below only "Direct-model-use" pose estimation approach corresponding to an explicit 3D geometric representation of human shape and kinematic structure by SQ.

Some authors propose recovering a pose with a shape detection stage (by hierarchical exemplar matching in the individual camera views with 3D upper body model based on tapered SQ), combining with Viterbi-style best trajectory estimation, and a filtering approach to 3D model texturing (Hofmann and Gavrilu, 2009). Other authors used a method for restoring 3D human body motion from monocular video sequences based on a robust image matching

metric, incorporation of joint limits and non-self-intersection constraints, and a sample-and-refine search guided by rescaled cost-function covariance (Sminchisescu and Triggs, 2003). There is also a method for recovering an object by SQ models with the recover-and-select paradigm, filling range images with a set of seeds (small SQ models), and increasing these seeds with a growth iteration approach selecting the suitable models. This approach was tried out on a wooden mannequin (Jaklic et al., 2000; Leonardis et al., 1997).

We propose using the hierarchical RANSAC-based model-fitting technique with a composite SQ model of human body and limbs. It is known that SQ models permit to describe complex-geometry objects with few parameters and generate simple minimization function to estimate an object pose (Jaklic et al., 2000 and Leonardis et al., 1997). We assume the body shape and dimensions are known a-priori to model body and limbs by SQ with correct anthropometric parameters in the metric coordinate system. The logic of our 3D Human Body pose estimation algorithm is presented by the block diagram (Figure 1). The object pose estimation starts with pre-processing of the 3D point cloud captured by multiple cameras. The preprocessing stage realizes segmentation of the Human Body into 9 parts (body, arms, forearms, hips and legs). After that the algorithm recovers 3D position of the body as the largest object ("Body Pose Search") and then

uses the information about body position to restore human limbs poses (“Limbs Pose Search”). To cope with measurement noise and outliers, the object pose is estimated by RANSAC-SQ-fitting technique. We control the fitting quality by setting inlier thresholds for limbs (body). These thresholds are a ratio of the optimal amount of inliers to whole data points of the corresponding limb (body). The tests showed that as a result of the Body Pose Search we can obtain a hypothesis with a slightly wrong body position, which can satisfy a body threshold, but can't be applied to overcome limb thresholds. For this reason, when the limb inliers solution less than a limb threshold, the algorithm restarts the Body Pose Search until finding suitable results of RANSAC-SQ-fitting for every limbs.

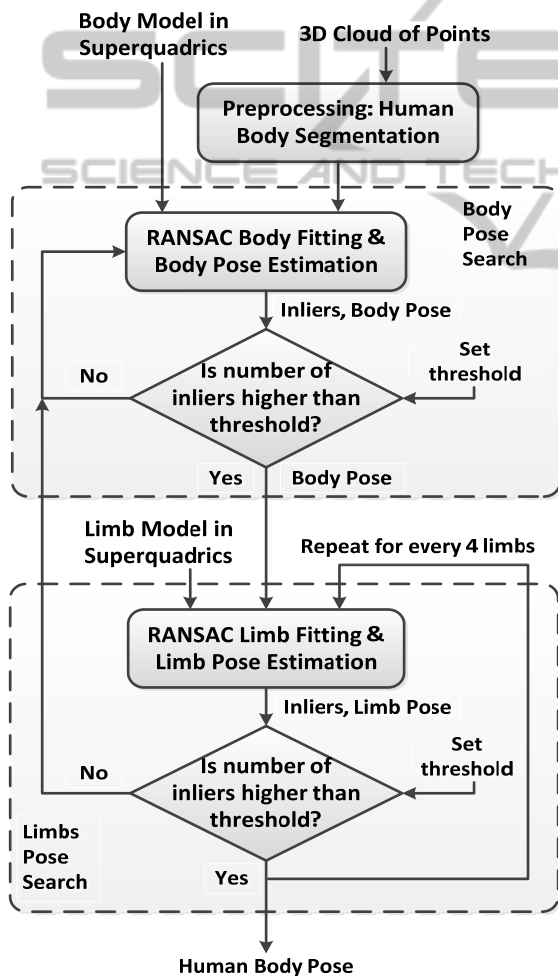


Figure 1: The block diagram of 3D Human Body Pose Estimation algorithm.

2 SUPERQUADRICS MODEL OF THE HUMAN BODY

2.1 SuperQuadric parameters

It is known (Jaklic et al., 2000 and Leonardis et al., 1997) that the explicit form of the parametric equation of the superquadrics, which is usually used for SQ representation and visualization, is:

$$\begin{bmatrix} x \\ y \\ z \end{bmatrix} = \begin{bmatrix} a_1 \cdot \text{signum}(\cos \eta) \cdot |\cos \eta|^{\varepsilon_1} \cdot \text{signum}(\cos \omega) \cdot |\cos \omega|^{\varepsilon_2} \\ a_2 \cdot \text{signum}(\cos \eta) \cdot |\cos \eta|^{\varepsilon_1} \cdot \text{signum}(\sin \omega) \cdot |\sin \omega|^{\varepsilon_2} \\ a_3 \cdot \text{signum}(\sin \eta) \cdot |\sin \eta|^{\varepsilon_1} \end{bmatrix} \quad (1)$$

where x, y, z – superquadric coordinate system;
 a_1, a_2, a_3 – scale parameters of the object;
 $\varepsilon_1, \varepsilon_2$ – object shape parameters;
 η, ω – spherical coordinates.

The implicit superquadric equation is more suitable for mathematical modeling to do fitting 3D data:

$$F(x, y, z) = \left(\left(\frac{x}{a_1} \right)^{\varepsilon_2} + \left(\frac{y}{a_2} \right)^{\varepsilon_2} \right)^{\frac{\varepsilon_1}{\varepsilon_2}} + \left(\frac{z}{a_3} \right)^{\frac{\varepsilon_1}{\varepsilon_2}} \quad (2)$$

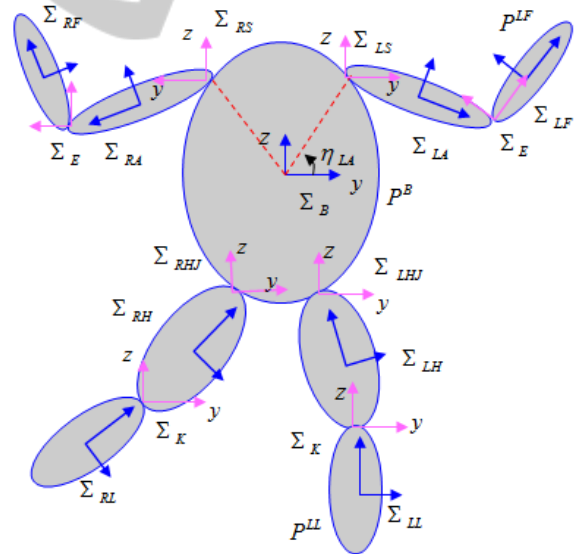


Figure 2: Presentation of Human Body in 9 parts: B – body, LA/RA – Left/Right Arms, LF/RF – Left/Right Forearms, LH/RH – Left/Right Hips, LL/RL – Left/Right Legs. Other abbreviations: LS – Left Shoulder, E – Elbow, η_{LA} – angle position of Left Shoulder, LHJ – Left Hip Joint, K – Knee, etc.

The object under investigation is the Human Body (Figure 2), which consists of 9 superquadrics –

superellipsoids with the shape parameters $\varepsilon_1 = \varepsilon_2 = 0.5$ and the following scaling parameters for different parts of the body:

- Body: $a_1 = 0.095, a_2 = 0.18, a_3 = 0.275$ (m).
- Arms: $a_1 = a_3 = 0.055, a_2 = 0.15$ (m).
- Forearms: $a_1 = a_3 = 0.045, a_2 = 0.13$ (m).
- Hips: $a_1 = a_2 = 0.075, a_3 = 0.2$ (m).
- Legs: $a_1 = a_2 = 0.05, a_3 = 0.185$ (m).

The scale parameters of SQ are presented in the metrical superquadric object-centered coordinate systems.

2.2 Human Body in SQ

The position of Human Body is defined by the following rotation & translation sequences of the Body Superquadrics:

1. Translation of center of BODY (x_c, y_c, z_c), along x, y, z -coordinates.
2. Rotation α among x (clockwise).
3. Rotation β among y (clockwise).
4. Rotation γ among z (clockwise).

The rotation matrix of BODY R_{BODY} is:

$$R_{BODY} = \begin{bmatrix} 1 & 0 & 0 & 0 \\ 0 & \cos(\alpha) & -\sin(\alpha) & 0 \\ 0 & \sin(\alpha) & \cos(\alpha) & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} \cos(\beta) & 0 & \sin(\beta) & 0 \\ 0 & 1 & 0 & 0 \\ -\sin(\beta) & 0 & \cos(\beta) & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} \cos(\gamma) & -\sin(\gamma) & 0 & 0 \\ \sin(\gamma) & \cos(\gamma) & 0 & 0 \\ 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix} \quad (3)$$

The transformation matrix T_{BODY} for the BODY is:

$$T_{BODY} = R_{BODY} \cdot \begin{bmatrix} 1 & 0 & 0 & x_c \\ 0 & 1 & 0 & y_c \\ 0 & 0 & 1 & z_c \\ 0 & 0 & 0 & 1 \end{bmatrix} \quad (4)$$

2.3 Human Arms and Forearms in SQ

Let's consider the transformation equations for Left Arm and Forearm.

The position of Left Shoulder according to the center of the body coordinate system (Figure 2) is estimated by SQ explicit equation (1):

$$P_S^B = P \left(\eta = \eta_{LA}, \omega = \frac{\pi}{2} \right) = \begin{bmatrix} 0 \\ a_2 \cdot \text{signum}(\cos \eta) \cdot |\cos \eta|^{\varepsilon_1} \\ a_3 \cdot \text{signum}(\sin \eta) \cdot |\sin \eta|^{\varepsilon_1} \end{bmatrix} \quad (5)$$

Taking into account (5), the transformation Body - Left Shoulder ($B-LS$) will be:

$$T_{LS}^B = \begin{bmatrix} 1 & 0 & 0 & \dots \\ 0 & 1 & 0 & P_S^B \\ 0 & 0 & 1 & \dots \\ 0 & 0 & 0 & 1 \end{bmatrix} \quad (6)$$

We can express the transformation: Left Shoulder - Left Arm ($LS-LA$) by the following

rotation & translation sequences:

1. Rotation α among x (clockwise).
2. Rotation β among z (anticlockwise).
3. Rotation γ among y (clockwise).
4. Translation of SQ center on distance a_2 along y .

$$T_{LA}^{LS} = T_{LA}^{LS}(\alpha = \alpha_{LA}, \beta = \beta_{LA}, \gamma = \gamma_{LA}) = R_{LA} \cdot \begin{bmatrix} 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & a_2 \\ 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix} \quad (7)$$

where R_{LA} is the rotation matrix of Left Arm:

$$R_{LA} = \begin{bmatrix} 1 & 0 & 0 & 0 \\ 0 & \cos(\alpha) & -\sin(\alpha) & 0 \\ 0 & \sin(\alpha) & \cos(\alpha) & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} \cos(\beta) & \sin(\beta) & 0 & 0 \\ -\sin(\beta) & \cos(\beta) & 0 & 0 \\ 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} \cos(\gamma) & 0 & \sin(\gamma) & 0 \\ 0 & 1 & 0 & 0 \\ -\sin(\gamma) & 0 & \cos(\gamma) & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix} \quad (8)$$

The transformation Left Arm - Elbow ($LA-E$) is

$$T_E^{LA} = \begin{bmatrix} 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & a_2 \\ 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix} \quad (9)$$

The transformation Elbow - Left Forearm ($E-LF$) is created by

1. Rotation δ_{LF} among x (clockwise).
2. Translation of SQ center on $-a_2$ along y .

$$T_{LF}^E = T_{LF}^E(\delta = \delta_{LF}) = \begin{bmatrix} 1 & 0 & 0 & 0 \\ 0 & \cos(\delta) & \sin(\delta) & -a_2 \\ 0 & -\sin(\delta) & \cos(\delta) & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix}^{-1} \quad (10)$$

Finally, taking into account equations (5)-(10), the full transformation for every point of system "Body - Left Forearm" ($B-LF$) can be calculated this way:

$$P^B = T_{LS}^B \cdot T_{LA}^{LS} \cdot T_E^{LA} \cdot T_{LF}^E \cdot P^{LF} \quad (11)$$

$$P^{LF} = (T_{LS}^B \cdot T_{LA}^{LS} \cdot T_E^{LA} \cdot T_{LF}^E)^{-1} \cdot P^B.$$

where P^B, P^{LF} - coordinates of Body and Left Forearm points correspondingly (Figure 2).

The main equations for Right Arm and Forearm are calculated the same way.

2.4 Human Hips and Legs in SQ

Analogically with previous equations (Section 2.3), the full transformation for every point of system "Body - Left Leg" ($B-LL$) is calculated this way:

$$P^B = T_{LHJ}^B \cdot T_{LH}^{LHJ} \cdot T_K^{LH} \cdot T_{LL}^K \cdot P^{LL} \quad (12)$$

$$P^{LL} = (T_{LHJ}^B \cdot T_{LH}^{LHJ} \cdot T_K^{LH} \cdot T_{LL}^K)^{-1} \cdot P^B.$$

where P^B, P^{LL} - coordinates of Body and Left Leg points respectively (Figure 2);

T – corresponding transformations (13)-(16).

The transformation Body – Left Hip Joint ($B-LHJ$) is absolutely the same as T_{LS}^B from equation (6), except using the angle η_{LL} in the equation (5) for calculation of the Left Hip position.

The transformation Left Hip Joint – Left Hip ($LHJ-LH$) uses other rotation sequences and translation if compare with equations (7) and (8):

1. Rotation α among x (clockwise).
2. Rotation β among y (anticlockwise).
3. Rotation γ among z (clockwise).
4. Translation of SQ center on distance $-a_3$ along z .

$$T_{LH}^{LHJ} = T_{LH}^{LHJ}(\alpha = \alpha_{LH}, \beta = \beta_{LH}, \gamma = \gamma_{LH}) = R_{LH} \cdot \begin{bmatrix} 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 \\ 0 & 0 & 1 & -a_3 \\ 0 & 0 & 0 & 1 \end{bmatrix} \quad (13)$$

where R_{LH} is the rotation matrix of Left Hip:

$$R_{LH} = \begin{bmatrix} 1 & 0 & 0 & 0 \\ 0 & \cos(\alpha) & -\sin(\alpha) & 0 \\ 0 & \sin(\alpha) & \cos(\alpha) & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} \cos(\beta) & 0 & -\sin(\beta) & 0 \\ 0 & 1 & 0 & 0 \\ \sin(\beta) & 0 & \cos(\beta) & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} \cos(\gamma) & -\sin(\gamma) & 0 & 0 \\ \sin(\gamma) & \cos(\gamma) & 0 & 0 \\ 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix} \quad (14)$$

The transformation Left Hip - Knee ($LH-K$) is

$$T_K^{LH} = \begin{bmatrix} 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 \\ 0 & 0 & 1 & -a_3 \\ 0 & 0 & 0 & 1 \end{bmatrix} \quad (15)$$

The transformation Knee - Left Leg ($E-LL$) is created by

1. Rotation δ_{LL} among y (clockwise).
2. Translation of SQ center on a_3 along z .

$$T_{LL}^K = T_{LL}^K(\delta = \delta_{LL}) = \begin{bmatrix} \cos(\delta) & 0 & -\sin(\delta) & 0 \\ 0 & 1 & 0 & 0 \\ \sin(\delta) & 0 & \cos(\delta) & a_3 \\ 0 & 0 & 0 & 1 \end{bmatrix}^{-1} \quad (16)$$

The similar transformations for Right Hip and Leg are described by almost the same equations.

3 3D HUMAN BODY FITTING ALGORITHM

3.1 About Sensors, Object and Data

The 3D point cloud is captured with a multi-camera system developed at the University of Trento in the framework of the project VERITAS (De Cecco, Paludet, et al., 2010). The multi-camera system for

acquiring range images consists of 8 pairs of cameras, which are a multiple stereo system, like a multi-camera system described in the paper (De Cecco, Pertile, et al., 2010), employing measurements a 3D-surface with superimposed colored markers.

The multi-camera system gives 3D video of Human Body movement consisted of 119 frames, but we are analyzing every frame separately. The total amount of 3D Human Body data points for single 3D video frame is more than 2100.

3.2 Preprocessing: Segmentation

The segmentation of 3D point cloud of a human body has been done automatically basing on the clothing analysis. We extract the human being clusters (body and eight limbs: left/right arms, forearms, hips and legs) according to the special clothing marks on the garment (Figure 6). These marks generate color structures, which are pre-defined clothing models. The result of this clothing analysis is a segmentation matrix, the elements of which set belonging to the definite limbs of the body for every data points. Experimental results show that such clothing segmentation is well-able to extract limbs of human body from range images with variations in background environment and lighting conditions.

The segmentation is completed with the use of RANSAC fitting. The markers near to the body joints have uncertain association with limbs. This uncertainty can be solved using RANSAC-SQ-fitting. As the result of this clothing segmentation we have approximately 800 data points of the body, 30-70 points of left/right arms, 15-25 points of forearms, 300-600 points of hips, and 80-150 points of legs (Figures 6 and 7).

This method will also work with any 3D point cloud data acquired by other sensors (for example Kinect) with following segmentation of body parts from single depth images invariantly to body clothing, as an example, using randomized decision forests (Shotton J., et al., 2011).

3.3 RANSAC Algorithm

We use RANSAC ("RANDOM SAmple Consensus") algorithm in estimating 3D Body and Limb Poses with SQ Model Fitting. To remind the basic concept of RANSAC algorithm, Figure 3 presents the pseudocode of RANSAC algorithm based on Peter Kovesi software (Kovesi, 2008). The number of iterations performed by RANSAC (the parameter k) can be determined from the following formula:

$$k = \frac{\log(1-p)}{\log\left(1 - \left(\frac{\text{inliers}}{n}\right)^s\right)} \quad (17)$$

where p – is the probability desired for choosing at least one sample free from outliers (in most of applications: $p=0.99$);

s – is a number of points required to fit the model.

The success of RANSAC usage depends on choosing the right models. In our case it means correct choice of SQ parameters (as anthropometric sizes of human body and limbs) and logic of recognition (i.e. the sequence of body/limbs fitting). The attempt of one stage RANSAC recognition whole parts of human body simultaneously will be failed because of big amount of outliers. The test showed that an acceptable quality of RANSAC-SQ-fitting can be achievable by hierarchical human body pose estimation (Figure 1).

3.4 RANSAC Model Fitting

The Body and Limbs Pose Search stages of the algorithm are very similar and have common logic and functions (Figure 1). The logic of RANSAC-SQ-fitting algorithms both for the body and a limb are explained by pseudocodes (Figures 4 and 5).

Let's consider the RANSAC Body Fitting algorithm. We are using the RANSAC-based Object Search to find the body pose hypothesis, i.e. 6 variables: 3 angles of rotation (α, β, γ) and 3 translation coordinates (x_C, y_C, z_C). Having these variables we can calculate the transformation matrix T_{BODY} (4). We are fitting a model described by the superquadric implicit equation (2) to 3D data of the known object (i.e. the points of the body sorted by segmentation). Each RANSAC sample calculation is started by picking a set of random points ($s = 6$ points for Body fitting, which are the minimal number of points to calculate the SQ position) from 3D datapoints in the world coordinate system (x_{wi}, y_{wi}, z_{wi}). To transform these points to the superquadric centered coordinate system (x_{si}, y_{si}, z_{si}), we use the following equation:

$$F_{si}(x_{si}, y_{si}, z_{si}) = T_{BODY}^{-1} \cdot \begin{bmatrix} x_{wi} \\ y_{wi} \\ z_{wi} \\ 1 \end{bmatrix} \quad (18)$$

where T_{BODY}^{-1} is the inverting homogeneous transformation matrix of the body (4).

Then we are calculating the inside-outside

function according to the superquadric implicit equation (2) in world coordinate system:

$$F_{wi} = \left[\left(\frac{F_s(x_{si})}{a_1} \right)^{\frac{2}{\epsilon_2}} + \left(\frac{F_s(y_{si})}{a_2} \right)^{\frac{2}{\epsilon_2}} \right]^{\frac{\epsilon_2}{\epsilon_1}} + \left(\frac{F_s(z_{si})}{a_3} \right)^{\frac{2}{\epsilon_1}} \quad (19)$$

It is easy to see that the inside-outside function for superquadrics has 11 parameters (Jaklic et al., 2000; Solina, 1990):

$$F_{wi} = F(x_{wi}, y_{wi}, z_{wi}, a_1, a_2, a_3, \epsilon_1, \epsilon_2, \alpha, \beta, \gamma, x_C, y_C, z_C), \quad (20)$$

where 5 parameters of the superquadric size and shape are known ($a_1, a_2, a_3, \epsilon_1, \epsilon_2$) and other 6 parameters ($\alpha, \beta, \gamma, x_C, y_C, z_C$) represent the orientation and position of superquadrics in space and should be found by minimizing the cost-function:

$$\min_{F_w} \sum_{i=1}^s F_i(x_i)^2 = \left(F_{wi}^{\epsilon_1} - 1 \right)^2, \quad (21)$$

where additional exponent ϵ_1 ensures that the points of the same distance from SQ surface have the same values of F_w (Solina and Bajcsy, 1990).

So we are fitting SQ model to this random dataset by minimizing an inside-outside function of distance to SQ surface (applying the “Trust-Region algorithm” or “Levenberg-Marquardt algorithm” in the nonlinear least-square minimization method). After this we are evaluating amount of inliers by comparing the distances between every point of 3D point cloud and SQ model with assigned distance threshold t (to accelerate the calculations we took the distance threshold $t = 2 \text{ cm}$):

$$d_i = \sqrt{a_1 \cdot a_2 \cdot a_3 \cdot \left(F_{wi}^{\epsilon_1} - 1 \right)}. \quad (22)$$

Analogically, we are realizing the RANSAC Limb Fitting (Figure 5). Let's consider the example of the Limb Fitting of Left Arm (LA) and Forearm (LF). The main differences between RANSAC Body and Limb Fitting are:

- in the minimal number of points to calculate the SQ position for Limb fitting $s = 3$ (although we need to set the body transform matrix T_{BODY} , obtained from the Body Fitting algorithm).
- in using 4 variables for Limbs Pose Search: 4 angles of rotation ($\alpha, \beta, \gamma, \delta$) equations (8,10).
- in minimizing the joint cost-function of two superquadrics together, considering two limbs simultaneously:

$$\min_{F_w} \sum_{i=1}^s F_i(x_i)^2 = \left[\left(F_{wi(LA)}^{\epsilon_1^{LA}} - 1 \right) \cdot \left(F_{wi(LF)}^{\epsilon_1^{LF}} - 1 \right) \right]^2, \quad (23)$$

where abbreviations *LA* and *LF* mean that parameters and variables are related to the Left Arm (LA) and Forearm (LF) Limbs correspondingly.

```

Algorithm RANSAC (x, fittingfn,
                  distfn, s, t, Trials)

% x - a dataset  $x^n$  of n observations
% fittingfn - a function that fits a model to x
% distfn - a function that checks a distance
%           from a model to x
% s - min number of data to fit the model M
% t - a threshold (a distance: datapoint - model)
% Trials - a number of iterations in algorithm

iter := 0           % count of iterations
bestM := 0         % the best model
inliers := 0      % accumulator for inliers
score := 0        % amount of inliers
p := 0.99        % probability of a sample
                % without outliers

while k > iter
% randomly selected j values from data  $x^n$ 
 $x_j^s := \text{random}(x^n)$ ;
% model parameters, which fitted to  $x_j^s$ 
M := fittingfn( $x_j^s$ );
for all  $x_i$  from  $x^n$ 
    if distfn(M,  $x_i^n$ ) < t
        inlk :=  $x_i$ 
    end if
end for
% amount of inliers
m := length(inlm);
% the test to check how good the model is
if m > score
    score := m; % amount of inliers
    inliers := inl; % inliers
    bestM := M; % the best model

    
$$k = \frac{\log(1 - p)}{\log\left(1 - \left(\frac{\text{inliers}}{n}\right)^s\right)}$$

end if
    increment iter
if iter > Trials
    break
end if
end while

```

Figure 3: Pseudocode of RANSAC algorithm.

To speed up the fitting process, the position of initial starting point of minimization searching in world coordinates can be chosen in the center of gravity of the body. Thus SQ modeling allows to

recovery an object in “clouds of points” with using the limited number of 3D data points. The minimization process with “Trust-Region” or “Levenberg-Marquardt” algorithms is stable without

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Algorithm RANSAC_Body_Fitting (x, t)

s = 6;           % min number of points to fit a SQ
t = 2 · 10-2; % a threshold: datapoint-SQ (2 cm)
% Trials - a number of iterations in algorithm
% x - a dataset  $x^n$  of n points of a body, which
% are a vector of the world coordinates ( $x_{w_i}, y_{w_i}, z_{w_i}$ )
% fittingfn - function to define SQ position by s, x.
function fittingfn ( $x^s$ )
    set  $x_0^s = 0$ ; % initial values of  $\alpha, \beta, \gamma, x_c, y_c, z_c$ 
    set SQBODY parameters  $a_1 - a_3, \varepsilon_1, \varepsilon_2$ 
    for all  $x_i$  from  $x^n$ 
        
$$T_{BODY} = \begin{bmatrix} 1 & 0 & 0 & 0 \\ 0 & \cos(\alpha) & -\sin(\alpha) & 0 \\ 0 & \sin(\alpha) & \cos(\alpha) & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} \cos(\beta) & 0 & \sin(\beta) & 0 \\ 0 & 1 & 0 & 0 \\ -\sin(\beta) & 0 & \cos(\beta) & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} \cos(\gamma) & -\sin(\gamma) & 0 & 0 \\ \sin(\gamma) & \cos(\gamma) & 0 & 0 \\ 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} 1 & 0 & 0 & x_c \\ 0 & 1 & 0 & y_c \\ 0 & 0 & 1 & z_c \\ 0 & 0 & 0 & 1 \end{bmatrix}$$

        
$$F_s(x_i, y_i, z_i) = T_{BODY}^{-1} \cdot \begin{bmatrix} x_{w_i} \\ y_{w_i} \\ z_{w_i} \\ 1 \end{bmatrix};$$

        
$$F_w = \left( \left( \frac{F_s(x_i)}{a_1} \right)^{\frac{2}{\varepsilon_1}} + \left( \frac{F_s(y_i)}{a_2} \right)^{\frac{2}{\varepsilon_2}} \right)^{\frac{\varepsilon_2}{\varepsilon_1}} + \left( \frac{F_s(z_i)}{a_3} \right)^{\frac{2}{\varepsilon_1}};$$

        
$$\min_{F_w} \sum_{i=1}^s F_i(x_i)^2 = (F_w^{\varepsilon_1} - 1)^2.$$

    end for
    calculate variables  $\alpha, \beta, \gamma, x_c, y_c, z_c$ 
    by minimizing  $\min \sum_{i=1}^s F_i(x_i)^2$ 

return  $T_{BODY}$ 

% distfn - a function to select distances from SQ to x
function distfn ( $T_{BODY}, x$ )
    for all  $x_i$  from  $x^n$ 
        
$$F_s(x_i, y_i, z_i) = T_{BODY}^{-1} \cdot \begin{bmatrix} x_{w_i} \\ y_{w_i} \\ z_{w_i} \\ 1 \end{bmatrix};$$

        
$$F_w = \left( \left( \frac{F_s(x_i)}{a_1} \right)^{\frac{2}{\varepsilon_1}} + \left( \frac{F_s(y_i)}{a_2} \right)^{\frac{2}{\varepsilon_2}} \right)^{\frac{\varepsilon_2}{\varepsilon_1}} + \left( \frac{F_s(z_i)}{a_3} \right)^{\frac{2}{\varepsilon_1}};$$

        
$$d_i = \sqrt{a_1 \cdot a_2 \cdot a_3 \cdot (F_w^{\varepsilon_1} - 1)^2}.$$

        if  $d_i < t$  then  $x_i = \text{inliers}$ 
    end for
return inliers

Trials = 1000; % a number of iterations
start RANSAC (x, fittingfn, distfn, s, t, Trials)
return best $T_{BODY}$ , bestinliers

```

Figure 4: Pseudocode of RANSAC Body Fitting algorithm.

Algorithm RANSAC_Limb_Fitting (\mathbf{x}, t)

$s = 3$; % min number of points to fit a SQ
 $t = 2 \cdot 10^{-2}$; % a threshold: datapoint-SQ (2 cm)
set T_{BODY} ; % from Body Fitting Algorithm
 % \mathbf{x} - a dataset \mathbf{x}^n of n points of a limb, which are
 a vector of the world coordinates (x_w, y_w, z_w)

% fittingfn - function to define SQ position by s, x .
function fittingfn (x, T_{BODY})
set $x_0^s = 0$; % initial values of $\alpha, \beta, \gamma, \delta$
set $a_1 - a_3, \varepsilon_1, \varepsilon_2$ for SQ_{LA} and SQ_{LF}
set η_{LA} ; % the angle position of a Shoulder
for all \mathbf{x}_i from \mathbf{x}^n

$$T_{LMB}^{LA} = T_{BODY} \cdot T_{LS}^{BODY} \cdot T_{LA}^{LS}, \quad T_{LMB}^{LF} = T_{BODY} \cdot T_{LS}^{BODY} \cdot T_{LA}^{LS} \cdot T_E^{LE} \cdot T_{LF}^{LE},$$

$$T_{LS}^{BODY}(\eta_{LA}) = \begin{bmatrix} 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & a_1^s \cos(\eta_{LA})^{\varepsilon_1} \\ 0 & 0 & 1 & a_1^s \sin(\eta_{LA})^{\varepsilon_1} \\ 0 & 0 & 0 & 1 \end{bmatrix}; \quad T_{LA}^{LS} = R_{LA} \cdot \begin{bmatrix} 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 \\ 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix};$$

$$R_{LA} = \begin{bmatrix} 1 & 0 & 0 & 0 \\ 0 & \cos(\alpha) & -\sin(\alpha) & 0 \\ 0 & \sin(\alpha) & \cos(\alpha) & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix} \cdot \begin{bmatrix} \cos(\beta) & \sin(\beta) & 0 & 0 \\ -\sin(\beta) & \cos(\beta) & 0 & 0 \\ 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix} \cdot \begin{bmatrix} \cos(\gamma) & 0 & \sin(\gamma) & 0 \\ 0 & 1 & 0 & 0 \\ -\sin(\gamma) & 0 & \cos(\gamma) & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix}$$

$$T_E^{LE} = \begin{bmatrix} 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & d^{LA} \\ 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix}; \quad T_E^{LE} = T_E^{LE}(\delta = \delta_{LF}) = \begin{bmatrix} 1 & 0 & 0 & 0 \\ 0 & \cos(\delta) & \sin(\delta) & -d^{LF} \\ 0 & -\sin(\delta) & \cos(\delta) & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix}$$

$$F_{x_i}^{LA}(x_i, y_i, z_i) = (T_{LMB}^{LA})^{-1} \cdot \begin{bmatrix} x_i \\ y_i \\ z_i \\ 1 \end{bmatrix}; \quad F_{x_i}^{LF}(x_i, y_i, z_i) = (T_{LMB}^{LF})^{-1} \cdot \begin{bmatrix} x_i \\ y_i \\ z_i \\ 1 \end{bmatrix};$$

$$F_w = \left(\frac{F_i(x_i)}{a_1} \right)^{\frac{2}{\varepsilon_1}} + \left(\frac{F_i(y_i)}{a_2} \right)^{\frac{2}{\varepsilon_2}} + \left(\frac{F_i(z_i)}{a_3} \right)^{\frac{2}{\varepsilon_3}};$$

$$\min_{F_i} \sum_{i=1}^s F_i(x_i)^2 = \left[\left(F_{w_i(LA)}^{\varepsilon_1} - 1 \right) \cdot \left(F_{w_i(LF)}^{\varepsilon_2} - 1 \right) \right]^2.$$

end for
calculate variables $\alpha, \beta, \gamma, \delta$ by
 minimizing $\min_{i=1}^s F_i(x_i)^2$

return $\alpha, \beta, \gamma, \delta$

% distfn - a function to select distances from SQ to x
function distfn ($x, T_{BODY}, \alpha, \beta, \gamma, \delta$)
for all \mathbf{x}_i from \mathbf{x}^n

$$F_{x_i}^{LA}(x_i, y_i, z_i) = (T_{LMB}^{LA})^{-1} \cdot \begin{bmatrix} x_i \\ y_i \\ z_i \\ 1 \end{bmatrix}; \quad F_{x_i}^{LF}(x_i, y_i, z_i) = (T_{LMB}^{LF})^{-1} \cdot \begin{bmatrix} x_i \\ y_i \\ z_i \\ 1 \end{bmatrix};$$

$$F_w = \left(\frac{F_i(x_i)}{a_1} \right)^{\frac{2}{\varepsilon_1}} + \left(\frac{F_i(y_i)}{a_2} \right)^{\frac{2}{\varepsilon_2}} + \left(\frac{F_i(z_i)}{a_3} \right)^{\frac{2}{\varepsilon_3}};$$

$$d_i = \sqrt{a_1 \cdot a_2 \cdot a_3 \cdot (F_w^{\varepsilon_1} - 1)^2}.$$

if $d_i < t$ **then** $\mathbf{x}_i = \text{inliers}$
end for
return *inliers*

Figure 5: Pseudocode of RANSAC Limb Fitting algorithm on example of Left Arm (LA) and Forearm (LF) Limbs.

redundant complexity and time consuming. The Figures 6 and 7 show the result of fitting by the RANSAC-SQ-Fitting algorithm (pink points – inliers, cyan – outliers). For most of 3D video frames, the amount of inliers is more than 65% from approximately 2100 points of 3D rawdata.

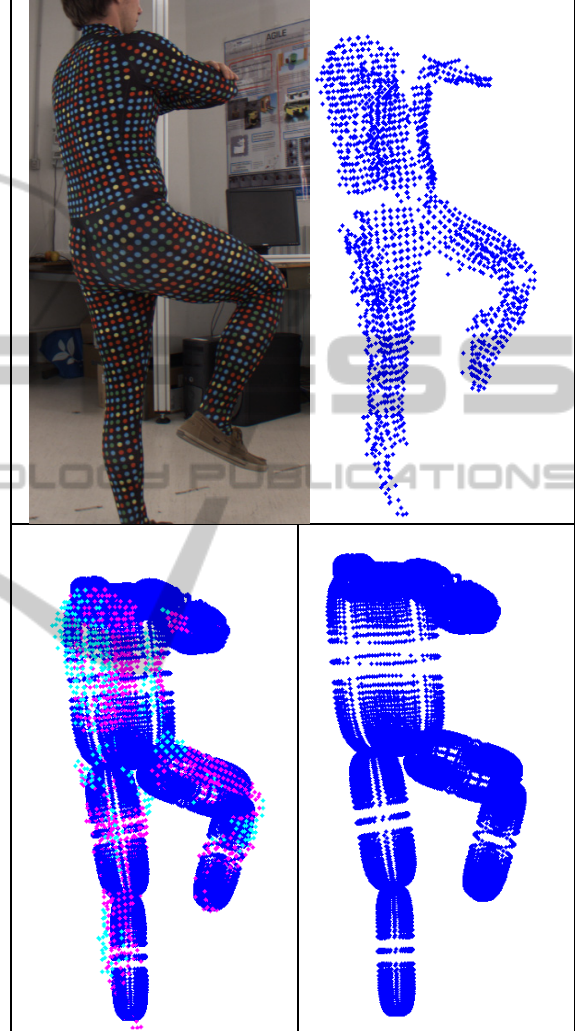


Figure 6: Illustration of RANSAC Limb Fitting algorithm. At the top: left – a pose of a human in the garment, right – “cloud of points”. At the bottom: left – the result of RANSAC-fitting to 3D data (pink points – inliers, cyan – outliers), right – final pose estimation.

The small amount of data points for arms and forearms (Section 3.2) gives some displacements of the upper limbs poses from one 3D video frame to another. It spoils the impression of the Human Body movement when we are preparing 3D video collecting together 3D Human body models from the individual video frames processed by RANSAC-SQ-fitting. This problem can be solved in future by

correcting 3D Human Body Pose Estimation algorithm or improving 3D data point acquisition process, or using other sensors (for example MS Kinect) and other segmentation techniques

4 RESULTS

The Figures 6 and 7 show the workability of the RANSAC-SQ-fitting algorithm for tasks of Human Body Pose Estimation. For most of 3D video frames, the amount of inliers is more than 65% from approximately 2100 points of 3D rawdata.

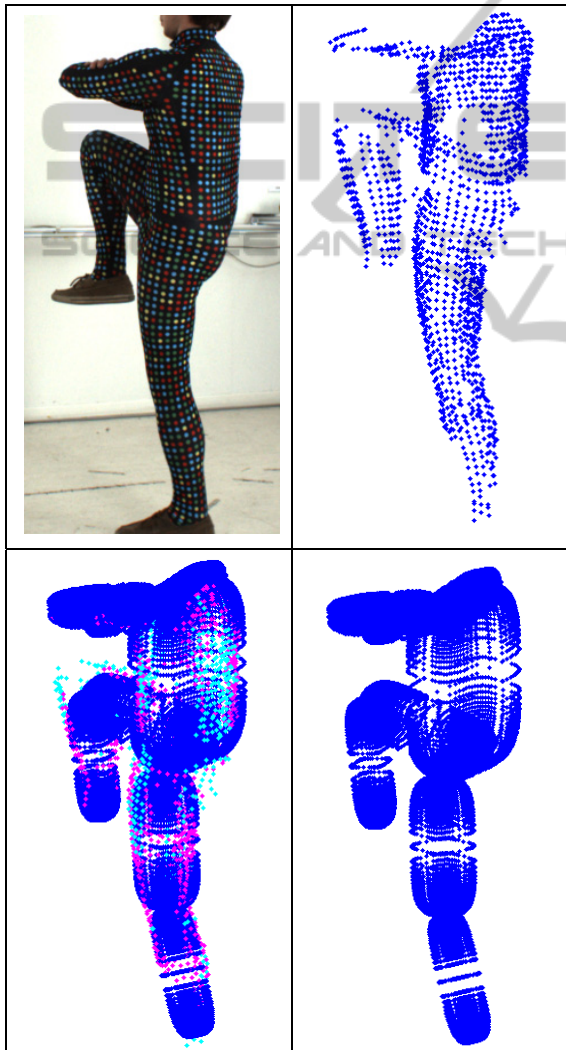


Figure 7: Illustration of RANSAC Limb Fitting algorithm. At the top: left – a pose of a human in the garment, right – “cloud of points”. At the bottom: left – the result of RANSAC-fitting to 3D data (pink points – inliers, cyan – outliers), right – final pose estimation.

The small amount of data points for the upper limbs gives some limb pose displacements and spoils the impression from body movement if collecting back 3D video from single frames.

This problem can be solved in future by correcting 3D Human Body Pose Estimation algorithm, or improving 3D data point acquisition process, or using other sensor and segmentation techniques.

The algorithm has been developed in MATLAB. 3D data have been captured from a multi-camera system and then processed offline. The pose estimation technique described has been tested at processing 3D video of Human Body movement consisted of 119 frames giving encouraging results. The presence of loops in the algorithm (during the hierarchical Body and Limbs Pose searches) can be a problem for real-time body movement application. But the correct comparative evaluation of speed and accuracy of the body pose estimation can be realized in the future if a) use other pose recognition methods with the existing multicamera system, or b) apply the proposed method with another sensor.

5 CONCLUSIONS

This paper describes a method of Human Body pose estimation from 3D real data obtained by a multi-camera system and structured by the special clothing analysis. This method will also work with any 3D point cloud data acquired by other sensors and segmented using any other algorithms.

The human body was modeled by a composite SuperQuadric (SQ) model presenting body and limbs with correct a-priori known anthropometric dimensions. The proposed method based on hierarchical RANSAC-object search with a robust least square fitting SQ model to 3D data: at first the body, then the limbs. The solution is verified by evaluating the matching score (the number of inliers corresponding to a-piori chosen distance threshold), and comparing this score with admissible inlier threshold for the body and limbs.

This method can be useful for applications dealt with 3D Human Body recognition, localization and pose estimation.

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