

Reconstructing 3D Pose and Motion from a Single Camera View

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Abstract

This paper presents a model based approach to human body tracking in which the 2D silhouette of a moving human and the corresponding 3D skeletal structure are encapsulated within a non-linear Point Distribution Model. This statistical model allows a direct mapping to be achieved between the external boundary of a human and the anatomical position. It is shown how this information, along with the position of landmark features such as the hands and head can be used to reconstruct information about the pose and structure of the human body from a monoscopic view of a scene.

1 Introduction

The human vision system is adept at recognising the position and pose of an object, even when presented with a monoscopic view. In situations with low lighting conditions in which only a silhouette is visible, it is still possible for a human to deduce the pose of an object. This is through structural knowledge of the human body and its articulation.

A similar internal model can be constructed mathematically which represents a human body and the possible ways in which it can deform. This information, encapsulated within a Point Distribution Model [5] can be used to locate and track a body. By introducing additional information to the PDM that relates to the anatomical structure of the body, a direct mapping between skeletal structure and projected shape can be achieved.

This work investigates the feasibility of such an approach to the reconstruction of 3D structure from a single view. To further aid the tracking and reconstruction process, additional information about the location of both the head and hands is combined into the model. This helps disambiguate the model and provides useful information for both its initialisation and tracking within the image plane.

1.1 Point Distribution Models

Point Distribution Models (PDMs) have proven themselves an invaluable tool in image processing. The *classic formulation* combines local edge feature detection and a model based approach to provide a fast, simple method of representing an object and how its structure can deform. Each pose of the object is described by a vector $\mathbf{x}_e = (x_1, y_1, \dots, x_N, y_N)$, representing a set of points specifying the object shape. A training set of vectors is

then assembled for a particular model class. The training set is then aligned (using translation, rotation and scaling) and the mean shape calculated. To represent the deviation within the shape of the training set, Principal Component Analysis (PCA) is performed on the deviation of the example vectors from the mean. This provides a compact mathematical model of how the shape deforms [5].

1.1.1 Linear Principal Component Analysis

Typically PDMs are generated by hand to ensure the correct identification of landmark points around a contour. Automated techniques are common place, but often require user intervention in the identification of landmark points to reduce the non-linearity that this introduces. Non-linear models when represented by the linear mathematics of PCA manifest themselves as un-robust models, which allow deformation not presented within the original dataset [2] [9] [10]. These problems become more acute when the move to 3D models is considered.

It has been proposed by Kotcheff and Taylor that non-linearity introduced during assembly of a training set could be eliminated by automatically assigning landmark points in order to minimise the non-linearity of the corresponding training cluster [12]. This can be estimated by analysing the size of the linear PDM that represents the training set. The more non-linear a proposed formulation of training set, the larger the PDM needed to encompass the deformation. This was demonstrated using a small test shape and scoring a particular assignment of landmark points according to the size of the training set (gained from analysis of the principal modes and the extent to which the model deforms along these modes, i.e. the eigenvalues of the covariance matrix[12]). This was formulated as a minimisation problem, using a genetic algorithm. The approach performed well however at a heavy computation cost [12].

As the move to larger, more complex, models or 3D models is considered, where dimensionality of the training set is high, this approach becomes unfeasible. A more generic solution is to use accurate non-linear representations and recent years has seen a wealth of publications on this subject (Discussed in the next section).

1.1.2 Non-linear Principal Component Analysis

Sozou *et al* first proposed using polynomial regression to fit high order polynomials to the non-linear axis of the training set [9]. Although this compensates for some of the curvature represented within the training set, it does not adequately compensate for high order non-linearity which manifests itself in the smaller modes of variation as high frequency oscillations of shape. In addition, the order of the polynomial to be used must be selected and the fitting process is time consuming.

Sozou *et al* further proposed modelling the non-linearity of the training set using a backpropagation neural network to perform non-linear principal component analysis [10]. This performs well, however the architecture of the network is application specific, also training times and the optimisation of network structure are time consuming.

Heap and Hogg suggested using a log polar mapping to remove non-linearity from the training set [7]. This allows a non-linear training set to be projected into a linear space where PCA can be used to represent deformation. The model is then projected back into the original space. Although a useful suggestion for applications where the only non-linearity is pivotal and represented in the paths of the landmark points in the original model, it does not provide a solution for the heavier non-linearity generated by other sources. What is

required is a means of modelling the non-linearity accurately, but with the simplicity and speed of the linear model.

Several researchers have proposed alternatives, which utilise non-linear approximations, estimating non-linearity through the combination of multiple smaller linear models [2][3][4][7]. These approaches have been shown to be powerful at modelling complex non-linearity in extremely high dimensional feature spaces [2]. It has also been proposed that analysis of the path taken by the training set through the non-linear feature space can be used to provide a probabilistic model of how the model varies through shape space [7].

2 Building a Combined Non-linear Point Distribution Model for a Human

The point distribution model is constructed from three components: the position of the head and hands within the image frame; the 2D contour which represents the shape of the body silhouette; and the 3D structure of the body (see Fig 1). Each of these is generated separately from the training image sequence and then concatenated to provide a training vector representing all these attributes. The relative position of the head and hands is represented as the location of these features in the image frame. When concatenated this generates a six dimensional feature vector $V_H=(x_1,y_1,...x_3,y_3)$. The body contour, once extracted from the image, is resampled to a list of 400 connected points. These are concatenated into an 800 dimensional feature vector $V_C=(x_1,y_1,...x_{400},y_{400})$. Lastly the skeletal structure of the 3D model is represented by 10 3D points which produce a 30 dimensional feature vector V_S . The relative location of the hands and head helps to disambiguate the contour during tracking. It can also be used to estimate an initial location and shape for the body contour.

When combining information for statistical analysis via PCA it is important that constituent features ($V_H V_C V_S$) are scaled to ensure that any particular feature does not dominate the principal axis'. This can be done by minimising the eigen entropy as proposed by Sumpter, Boyle and Tillett [11]. However as all three components exist within the same co-ordinate frame and are directly linked, and thus such scaling is unnecessary.

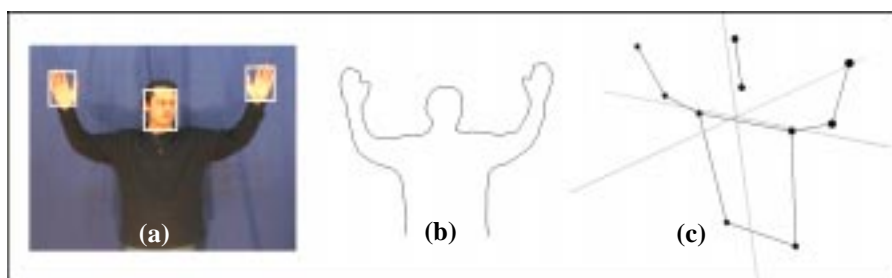


Figure 1 (a) Position of head and hands V_H (b) Body Contour V_C (c) Corresponding 3D model V_S

2.1 Hand and Head Position Estimation

Colour information is an important attribute within an image, which is often discarded.

McKenna, Gong and Raja have demonstrated that in a Hue-Saturation space, human skin occupies a relatively small cluster and can be used to segment a human head from a complex noisy scene. Using a gaussian mixture model to represent this colour space they have shown how multiple models for individuals can be used to probabilistically label an image and determine the most likely person present. Azarbayejani and Pentland have used similar methods to automatically segment both the hands and head from stereo image pairs, and using this, calculate their position and trajectories in 3D space [1].

By taking example images of human skin and plotting the constituent points in hue saturation space, a single gaussian cluster is sufficient to provide a reliable probabilistic measure for location of skin clusters in the image frame. By performing PCA on this colour cluster approximate bounds are calculated. If a sample pixel from a new image is within the Hue-Saturation bounds calculated from the PCA of skin samples then that pixel is marked as a probable location, producing a binary image. By performing erosion then dilation, noisy points are removed and clusters of marked skin points consolidated to blobs. A simple blobbing algorithm is then used to calculate approximate locations of skin artefacts within the image.



Figure 2. Blobs of skin artefacts

Figure 2 shows a sample image frame after processing. The results from the blobbing algorithm are used to calculate the centre and approximate size of the skin artefacts. This is used to place a cross over the segmented features for demonstration purposes.

This procedure is repeated for each image in the training sequence to extract the trajectories of the head and hand as the human moves.

2.2 Shape Extraction

For the purpose of simple contour extraction from the training set, shape extraction is facilitated through the use of a blue screen and chroma keying. This allows the background to be simply keyed out to produce a binary image of the body silhouette. As the figure always intersects the base of the image at the torso, an initial contour point is easily located. Once found, this is used as the starting point for a simple contour tracing algorithm which follows the external boundary of the silhouette and stores this contour as a list of connected points. In order to perform any statistical analysis on the contour, it must first be resampled to a fixed length. To ensure some consistency throughout the training set, landmark points are set at the beginning and end of the contour. A further landmark point is allocated at the highest point along the contour within 10 degrees of a vertical line drawn from the centroid

of the shape. Two further points are positioned at the leftmost and rightmost points of the contour. This simple landmark point identification results in non-linearity within the model. The problems associated with this are discussed in section 2.6.

2.3 Introducing 3D Information

The 3D skeletal structure of the human is generated manually. Co-ordinates in the xy (image) plane are derived directly from the image sequence by hand labelling. The position in the 3D dimension is then estimated for each key frame.

2.4 The Linear PDM

Once these separate feature vectors are assembled, they are concatenated to form an 836 dimensional vector which represents the total pose of the model. A training set of these vectors is assembled which represents the likely movement of the model. Figure 3 shows a sample of training images along with the corresponding contour and skeletal models in 2D.

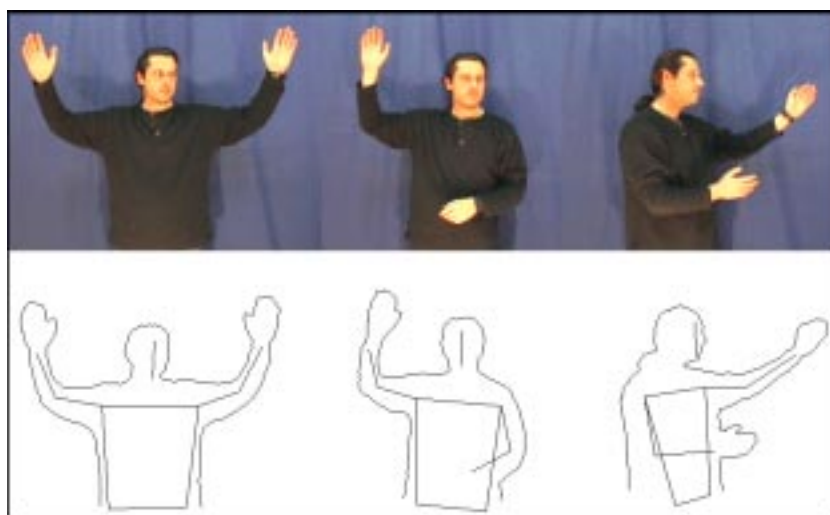


Figure 3.

Sample training images and corresponding contour and skeletal models

A linear PDM is now constructed [5] from the training set and its primary modes of variation are shown in Figure 4.

Figure 4 demonstrates the deformation of the composite PDM. The crosses are the locations of the hands and head. It can be seen that although the movement of the three elements are closely related, the model does not accurately represent the natural deformation of the body. The shapes generated by the primary modes of variation are not indicative of the training set due to its inherent non-linearity. In order to produce a model that is accurate/robust enough for practical applications, a more constrained representation is required.

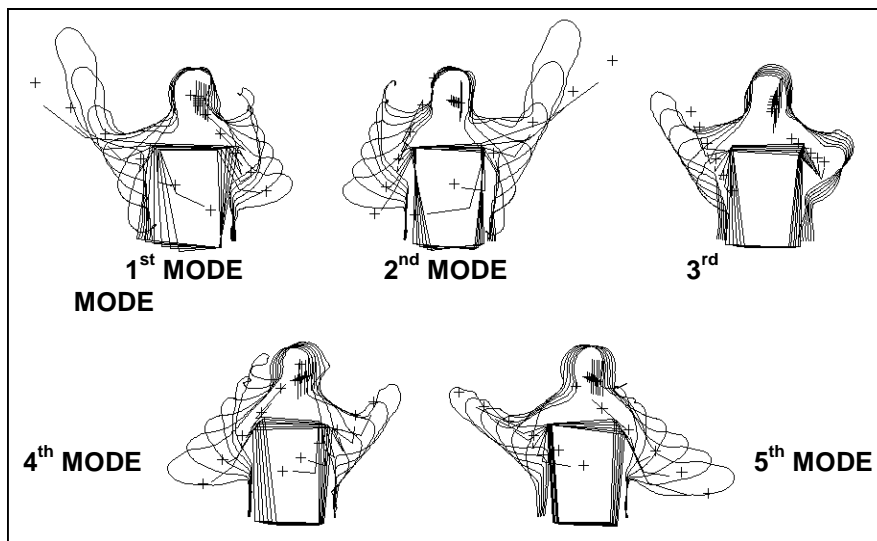


Figure 4, Primary modes of variation on the linear PDM

2.6 Non-Linear Estimation

Any analysis performed upon this training set is time consuming due to its high dimensionality and size. This dimensionality is therefore reduced by projecting the dataset down into a lower dimensional space while preserving the important information: its shape. The linear PDM model, which is unsuitable for modelling the deformation, is invaluable in this sense. Although the Linear PDM contains uncharacteristic deformation uncharacteristic to a human, it is capable of representing the original deformation contained in the training set [2]. After PCA, it is calculated that the first 84 eigenvectors that corresponding to the 84 largest eigenvalues encompass 99.99% of the deformation contained in the training set. By projecting each of the sample shapes down onto these vectors and recording their distance from the mean a new vector $R_V=(e_1, \dots, e_{84})$ is constructed. From this vector it is a simple back projection to reconstruct the model. Upon visual observation of the original vector and the reconstructed vector it can be seen that only the first 40 eigenvectors are necessary to represent the shape accurately. These primary 40 modes of deformation encompass 99.8% of the deformation. Projecting the entire training set down into this lower dimensional space achieves a dimensional reduction of 836 to 40, which significantly reduces the computation time required for further analysis.

In this lower dimensional space the information about the shape of the training set and how the model moves throughout it is preserved allowing further statistical analysis. Using a kmeans-clustering algorithm the space can be segregated into sub areas, which estimate the non-linearity [2]. Using standard cluster analysis the natural number of clusters can be estimated to be 25. By performing further PCA on each of the 25 clusters, the shape of the model can be constrained by restricting the shape vector to remain within this volume.

Figure 5 shows the training set after dimensional reduction from the original linear PDM, projected into 2 dimensions. The bounding boxes represent the 25 clusters which best estimate the curvature. These bounding boxes are the bounds of the first and second

modes of deformation for each of the sub PCA models. The number of modes for each cluster varies according to the complexity of the training set at that point within the space. All clusters are modelled to encompass 99.9% of the deformation within that cluster.

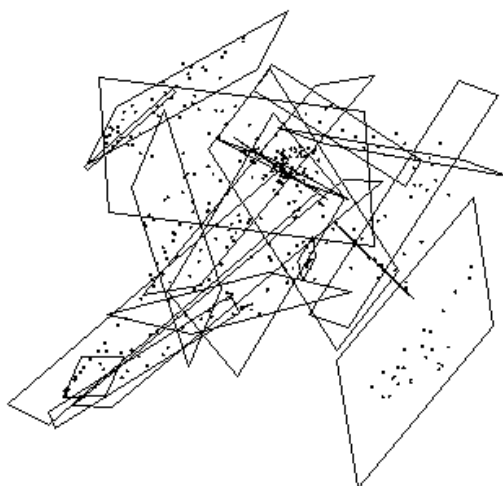


Figure 5 - Clusters in reduced shape space

3 Applying the PDM to an Image

3.1 Initialising the PDM

Upon initialisation the first step is to locate the position of the head and hands. This can be done via the procedure described in section 2.1 which does not need to be repeated on every iteration. Once done these positions can be used to initialise the PDM and give an initial guess as to the shape of the contour to be found. As it is not clear which blobs correspond to which features, three possible contours can be produced. The contour that iterates to the best solution provides the final state from which tracking proceeds.

3.2 Tracking with the PDM

Once initialised the two components must be fitted to the image separately. The contour is attracted to high intensity gradients within the image using local edge detection. The hand and head positions are used as centres in a single iteration of a kmeans-clustering algorithm on the segmented binary skin image. This is possible due to the assumption that the model will not change significantly from the last image frame.

3.3 Reconstruction of 3D Shape and Pose

As the shape deforms to fit with the image so the third element of the model, the skeleton, also deforms. By plotting this 3D skeleton, its movements mimic the motion of the human

in the image frame.

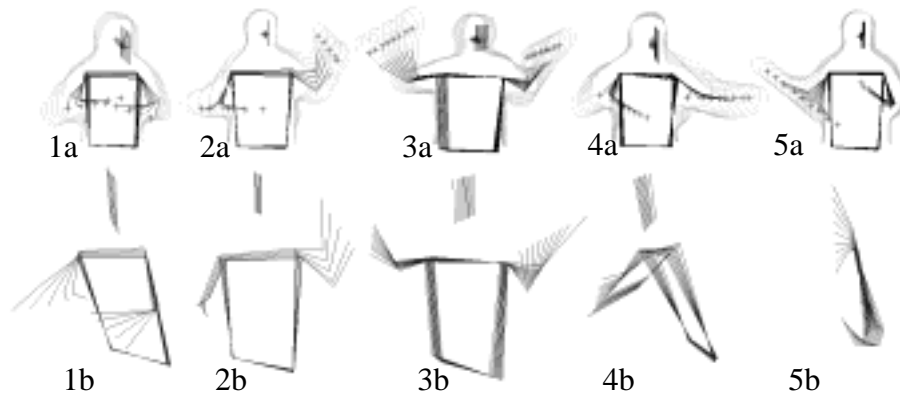


Figure 6 How the model deforms

Figure 6 demonstrates the correspondence between the body contour and skeletal structure. Each contour image (a) is generated from a different sub cluster of shape space. The deformation corresponds to the largest mode of deformation for that cluster. The 3D skeletal diagrams (b) correspond to the relevant contour (a), and demonstrate the movement of the skeleton. The orientation of these skeletal models has been changed in order to better visualise the movement in 3D. Skeleton (1b) demonstrates the arms moving in the z direction corresponding to the change in contour (1a) around the elbow region. Contour (4a) represents a body leant toward the camera with moving arms. Skeleton 4b shows the corresponding change in the skeleton with the shoulders twisting as the arms move. The Skeleton 5b is a plan view showing the movement of the hands.

All model points move along straight lines due to the linear clusters used to approximate the non-linear shape space. However, all poses of the models are lifelike human silhouettes, demonstrating the cluster PCAs ability at modelling the non-linearity.

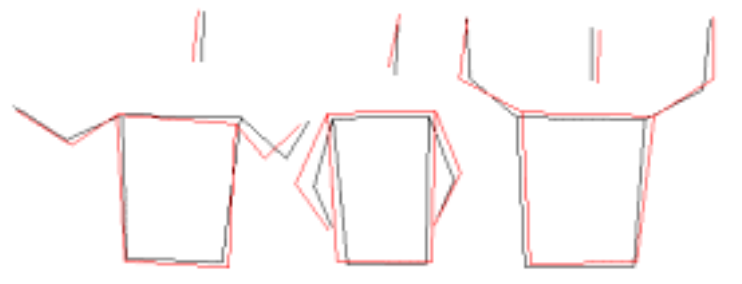


Figure 7 – Reconstructed poses from the model

Figure 7 shows the original model pose from the training set in grey with the reconstructed skeletal model in black. It can be seen that the original and reconstructed models are similar in pose and position with the length of limbs preserved, further demonstrating the absence of non-linear effects. However, as the constraints on shape space

are increased, so the performance degrades. Inconsistencies in the original and reconstructed models and the deterioration under heavy constraints can be attributed to the hand labelling of the training set. During hand labelling it is impossible to provide consistent models of the skeletal structure throughout the training set. This factor leads to the final model producing mean skeletal shapes which have been 'learnt' from the original training set and hence produces the inconsistencies observed in figure 7.

4 Conclusions

This paper has demonstrated how the 3D structure of an object can be reconstructed from a single view of its outline, using an internal model of shape and movement. The technique uses computationally inexpensive techniques for real time tracking and reconstruction of objects. It has also been shown how two sources of information can be combined to provide a direct mapping between them. The model appears to reconstruct 3D pose accurately. However, due to the acquisition of skeletal position, no ground truth information is available to test its accuracy. Being able to reconstruct 3D pose from a simple contour has applications in surveillance, virtual reality and smart room technology and could possibly provide an inexpensive solution to more complex motion capture modalities such as electromagnetic sensors and marker based vision systems.

5 Future Work

A full body model needs to be constructed for the generic tracking and pose estimation of humans. During its construction accurate skeletal information must be acquired to ensure usability of the resulting model. Key point trajectories will be acquired using electromagnetic sensors to provide training information during image acquisition for contour extraction. This will also provide the necessary ground truth data required to assess the accuracy of the final model. It is also possible to implement a stereoscopic or quad camera system to further increase accuracy and further disambiguate the contour fitting process. Inclusion of information about the orientation of the skeleton added during training would allow the estimation of contour shape in other image frames from a multi-camera system.

Acknowledgements

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