

11 **1. Introduction**

12 Human pose estimation [1, 1, 2] is widely applied in human-computer in-
13 teraction, smart video surveillance, health care, etc. Although a lot of efforts
14 have been devoted to the research of pose estimation, it remains a very chal-
15 lenging problem in computer vision because of occlusion, high dimensionality
16 of the search space and high variability in people’s appearance.

17 The depth image obtained by the depth sensor [3, 4, 5] can provide 2.5D
18 scene geometry, which facilitates both the segmentation of human body from
19 background and the disambiguation of similar poses. Recently, the focus of
20 pose estimation [6, 7, 8, 9] has been shifted toward pose estimation on depth
21 images. Most of these works can be divided into two categories: generative
22 methods and discriminative methods.

23 Typical generative methods include the proposals in [10, 9, 11, 12], in
24 which a kinematic chain and a 3D surface mesh are built as the human body
25 model. They treat the depth image as a point cloud over 3D space and apply
26 a model-fitting algorithm, such as the iterative closest point (ICP), to the
27 human body model to fit the 3D point cloud. Ye et al. [11], Ganapathi et
28 al. [12] and Baak et al. [9] combine dataset searching and model fitting to
29 approach the problem of 3D pose estimation. Ganapathi et al. [10] extend
30 the ICP to an articulated model by enforcing constraints over the pose space.
31 Although such methods do not need a training step, they suffer many draw-
32 backs. For example, the accuracy depends on the surface mesh level [13] and
33 the fitting usually needs long processing and inconvenient setups.

34 Compared with the generative methods, the discriminative methods do
35 not iteratively fit models to the observed data. Rather they directly esti-
36 mate the parameters about pose. Thus they can estimate the pose quickly
37 and adapt to various conditions. They regard the human pose as a collection
38 of different parts/joints and learn discriminative classifiers for the part/joint
39 detection [6, 8, 7, 14]. The most famous works on depth images are those
40 based on random forest [6, 8, 7]. Shotton et al. [6] formulate the pose estima-
41 tion as a classification task and use the random forests to learn the classifiers.
42 Girshick et al. [8] convert the classification task to the regression problem for
43 the estimation of the occluded parts. In [7], Sun et al. incorporate tem-
44 porary states of the object, such as person’s height and facing direction, to
45 boost the performance of the classifiers. However, these methods infer lo-
46 cations of body joints either independently [6, 8] or relying on some global
47 information [7], neglecting the dependence between body joints.

48 It is natural to boost the pose estimation performance by adding con-
49 straints among joints. One of the most widely used approach in this direc-
50 tion is to use graph model-based prior structure, which was first proposed
51 in [15] for general computer vision problems and later applied to the pose
52 estimation problem in [16]. It assumes that the relationships among joints
53 are state-constrained among the body parts. Two important components
54 are defined in the model: one is the appearance model which represents the
55 probability of a body part at a particular location in the given image; the
56 other is the prior model which represents the probability distribution over

57 pose space. To make a trade-off between computational efficiency and es-
58 timation accuracy, tree-structured models with a single Gaussian prior are
59 commonly used [15, 16, 17, 18]. However, as the diversity of human pose
60 increases, a simple Gaussian prior usually leads to a poor model of human
61 articulation, which cannot be applied well to the tasks on the depth images.
62 This is mainly due to two reasons. One is that it is not an easy work to
63 find a proper kernel number for the Gaussian model in a large dataset. A
64 small number may cause a poor fitting of the prior, while a large number
65 will cost extra computation and is prone to over-fitting. The other is that
66 the method always applies the same prior model to test samples, even when
67 they are of distinct poses. This limits the adaptability of the method. The
68 works in [19, 20] cluster poses into sub-clusters and learn a GMM for each
69 sub-cluster to enhance the adaptability of prior model. However, at the in-
70 ference stage, they need to infer all possible poses and select one as the final
71 output. This makes the inference complex.

72 In this paper, we propose a novel framework called Latent Variable Picto-
73 rial Structure (LVPS) for pose estimation on depth images. We construct and
74 estimate a latent variable based on the human silhouette. At the inference
75 stage, our model rebuilds the appearance model and the prior model based
76 on the values of the latent variable and then infers human poses. We shall
77 show its effectiveness through experiments on public datasets. Compared
78 with the state-of-the-art methods, our proposal can significantly increase the
79 accuracy of pose estimation.

80 The rest of the paper is organized as follows. We overview the proposal
81 in Section 2. Our LVPS model is introduced in Section 3 and its application
82 to the pose estimation in Section 4. We present experiments and discussions
83 in Section 5 and draw conclusions in Section 6.

84 2. Overview of the proposed method

85 Fig. 1 shows the framework of our LVPS. It consists of two main processes:
86 the training stage indicated by green arrows and the inference stage indicated
87 by blue arrows.

88 **The training stage.** The keys of the training stage involve generation
89 and selection of the latent variable and the training of models. In our work,
90 we extract silhouette features of poses, obtain their distributions, quantize
91 the distributions into a set of states C , and use the state label as the latent
92 variable. According to the value of the latent variable, all the training sam-
93 ples are partitioned into subsets. After that, we attach the value of the latent
94 variable to each sample and treat each sample as a two-labels object: a body
95 part label and a latent variable state. Samples with labels are then input
96 into classifiers to learn appearance models and prior models. As a result, the
97 diversity of the appearance and prior in each cluster would be reduced and
98 the prior model can be better learned and the discrimination ability of the
99 appearance model can be largely enhanced.

100 **The inference stage.** As the blue arrows indicate, to estimate one body
101 pose on depth image I , we shall first evaluate its latent state. This is, the

102 likelihood $p(c_i|I)$ is estimated. After that we rebuild our prior model and
 103 appearance model by assembling the learned models of individual clusters
 104 according to the likelihoods. As a result, our proposal adapts the models
 based on the specific test image.

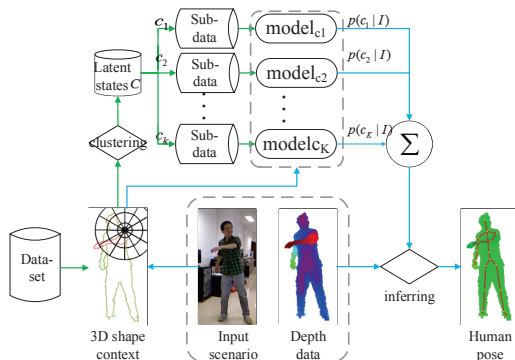


Figure 1: The flowchart of the proposed method: the process with green arrows is the training stage and that with blue arrows is the inference stage.

105

106 3. Latent variable pictorial structure

107 A classical pictorial structure model of the human body was proposed
 108 in [15]. It assumes that the dependences between body joints can be ex-
 109 pressed by a predefined graph, $G = (V, E)$, as shown in Fig 2, where V
 110 and E denote the sets of nodes and edges in the graph G , respectively. We
 111 use $X = \{x_1, x_2, \dots\}$ to denote the pose, in which x_i denotes the position of
 112 joint i . For the detection of an articular object, the objective function to be

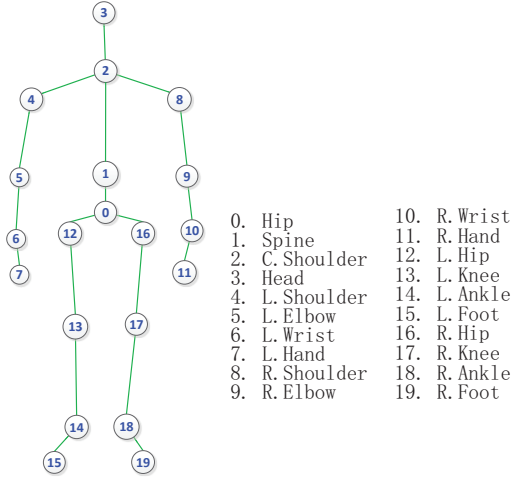


Figure 2: The graph model on human pose. The circle with a number is a vertex in V , which presents a joint/part of the body; the line between two joints is an edge in E , which indicates that the connected joints/parts are dependent.

113 maximized when given image I can be written as

$$p^{PS}(X|I) \propto \left\{ \prod_{i \in V} \phi(x_i|I) \right\} \left\{ \prod_{(i,j) \in E} \phi(x_i, x_j) \right\}, \quad (1)$$

114 where $\phi(x_i|I)$ denotes the appearance likelihood, which models the probabil-
 115 ity of a part at a particular location and orientation given the input image
 116 I , and the factor $\phi(x_i, x_j)$ denotes a prior, which models the probability dis-
 117 tribution over pose space. In this paper, the factor $\phi(x_i, x_j)$ describes the
 118 distribution of relative position between joint i and joint j .

119 In most existing methods based on the general pictorial structure model,
 120 only one tree-structured Gaussian prior is used to speed up the inference, and
 121 the appearance models of individual parts are learned independently. This

122 leads to a prior of low descriptive ability and an appearance model which
 123 cannot capture the multi-modal appearance of body parts, e.g. the different
 124 appearances of a body part in different views.

125 To overcome these issues, we incorporate a latent variable into the gen-
 126 eral pictorial structure to propose a latent variable pictorial structure model
 127 (LVPS). Specifically, we utilize the discrete state of the latent variable to par-
 128 tition samples and the pose space. Hence the diversity of the appearance and
 129 prior in each cluster would be reduced, which results in more effective and
 130 reliable appearance and prior models at the cluster level than the global mod-
 131 els. Besides, clustering over the latent variable feature space leads to a simple
 132 classifier. We use c to denote the discrete latent variable, $C = (c_1, \dots, c_K)$
 133 to denote the set of the K states of the latent variable, and $p(c_k|I)$ to denote
 134 the probability of the state c_k given image I .

135 Then based on the latent structure we obtain the posterior probability of
 136 pose X as

$$p^{LVPS}(X|I) \propto \sum_{c_k \in C} \{p^{PS}(X|c_k, I)p(c_k|I)\}, \quad (2)$$

137 where $p^{PS}(X|c_k, I)$ denotes the posterior probability conditional on the spe-
 138 cific cluster corresponding to c_k . The latent variable c may encode any de-
 139 sirable properties of the target objects. In this paper, we propose to utilize
 140 it to encode the whole human pose through body silhouette.

141 The inference stage is show in Fig 3. To the given image I , we first extract
 142 its latent variable value $Hist(I)$, which has a form of histogram of silhouette

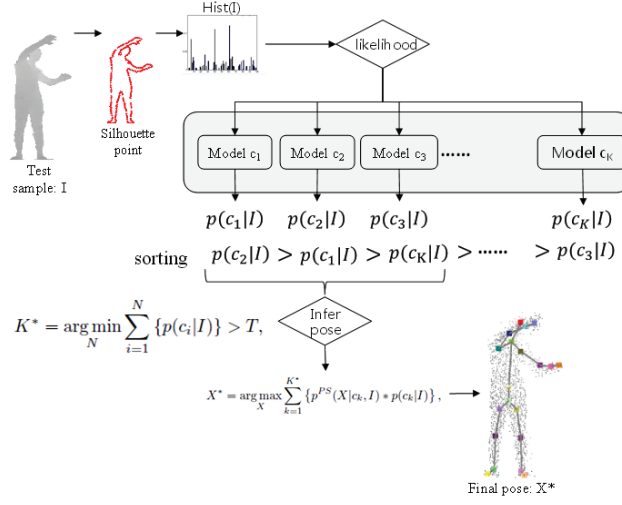


Figure 3: The flowchart of inferring a human pose X from the given image I .

143 features in this paper. Then the likelihood is evaluated between $Hist(I)$
 144 and those of sub-models. We sort the sub-models in descending order of the
 145 likelihood $p(c|I)$ and the first K^* sub-models that have their total likelihood
 146 beyond threshold T are selected. At last, a linear strategy is used to build
 147 the final detection model for the pose inference using (3):

$$\begin{aligned}
 X^* &= \arg \max_X \sum_{k=1}^{K^*} \{p^{PS}(X|c_k, I) * p(c_k|I)\}, \\
 K^* &= \arg \min_N \sum_{i=1}^N \{p(c_i|I)\} > T,
 \end{aligned} \tag{3}$$

148 where K^* is the number of selected sub-model, T is the threshold and N is
 149 a variable for counting sub-models. In this way we can adjust the number
 150 of the models used for the test sample, and its effect can be shown in the
 151 experiments in Section 5.4.

152 4. Details of LVPS

153 This section describes how the LVPS models are implemented for human
154 pose estimation. Since the samples are partitioned into subsets based on the
155 value of the latent variable, the variation of the pose space is decomposed
156 and a pose subspace can be better modeled even with a simple model. As a
157 result, two main parts will be discussed in the following: the generation and
158 selection of the latent variable and the learning of the appearance models.

159 4.1. The latent variable

160 A simple way to model the variation of the pose space is to cluster poses
161 directly in the pose space as in [20]. However, they have to learn a multinomial
162 logistic regression to classify each cluster. Another way is to use some
163 properties of the object [7, 21], such as torso orientation, person’s height or
164 facing direction. These features are natural, but they are not much associated
165 with the pose as a whole.

166 In our proposal, we extract a kind of silhouette feature to represent the
167 pose and use such feature to build our latent variable and to cluster our
168 samples. The most commonly used silhouette feature to represent a pose is
169 the shape context feature, which was first proposed in [22] for shape matching
170 and then used for human pose estimation [23, 24]. However, the silhouette
171 features extracted from RGB/grey images cannot represent the 3D structure
172 of pose. So He et al. [25] extend the 2D shape context [23] to 3D space.
173 To the best of our knowledge, existing pose estimation methods only use

174 silhouette feature to learn maps from the feature space to the pose space,
 175 rather than to build latent variables to boost the prior model.

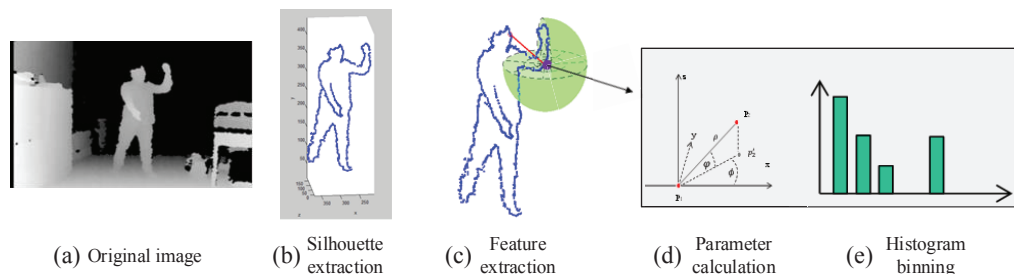


Figure 4: Extraction of shape context feature in [25]: (a) is the original depth image; (b) is the result after silhouette extraction; (c) shows how to extract the shape context feature on point p_1 ; (d) calculates the offset parameters between p_1 and any other points on the silhouette; (e) shows the building of histogram of the shape context feature.

176 In the following, we discuss how to generate and select the latent variable
 177 using the feature proposed in [25].

178 One brief flowchart of feature extraction [25] is shown in Fig 4. First,
 179 a sequence of edge points are extracted on each depth image. Then, for
 180 each edge point, the offsets between it and other points are calculated and
 181 voted into a histogram. This histogram encodes local pose information by
 182 collecting offsets on the edge points and is called shape context feature. At
 183 last, one pose is encoded by a bag of shape context features. More details
 184 about the feature extraction can be found in [25]. However, with such a bag
 185 of features, it is computationally consuming to compare two images.

186 To tackle this issue, we use the method in [23] to align these shape context
 187 features, which will be then used to form a feature vector for the construction
 188 of our latent variable. Specifically, we run k-means on the shape context

189 features from all the training samples to obtain B quantized centers. To
 190 represent one pose, we softly vote the shape context features of one image
 191 onto these learned centers with Gaussian weights. Finally, each pose on a
 192 depth image can be represented by a B -dimensional feature vector f . In the
 193 experiments, we set B to 100 as with [23]. Feature vectors of some samples
 are shown in Fig. 5.

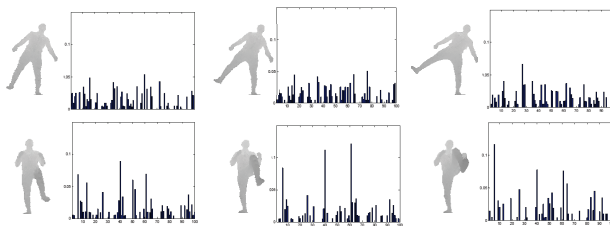


Figure 5: Some samples and their feature vectors f .

194

195 The feature vector f encodes the silhouette of body and can capture richer
 196 pose information than some straight properties, such as torso orientation and
 197 persons height. To quantize a feature vector further, we perform another k-
 198 means algorithm to obtain K discrete states (i.e. cluster labels) as the values
 199 of the feature vectors. We adopt the cluster label as the latent variable. After
 200 that, we can partition the training data into K subsets based on the value
 201 of the latent variable and can estimate the likelihood that image I belongs
 202 to the k th cluster c_k by using a simple histogram distance as

$$p(c_k|I) \propto 1/dst(Hist(I), Hist(c_k)), \quad (4)$$

203 where $dst(Hist(I), Hist(c_k))$ indicates the distance between two histograms

204 $Hist(I)$ and $Hist(c_k)$.

205 We show some average poses of individual clusters in Fig. 6. From Fig. 6,
206 we can find that by clustering poses through the mid-level representation we
207 can encode pose states and reduce the pose diversity in each cluster. For
208 example, in Fig. 6, clusters (1), (4) and (6) show the hands changes while
209 (1), (2) and (5) focus on the facing direction. In Section 5, more samples
210 are shown in the experiments and the value of sub-model number K and its
influence on the performance will be discussed.

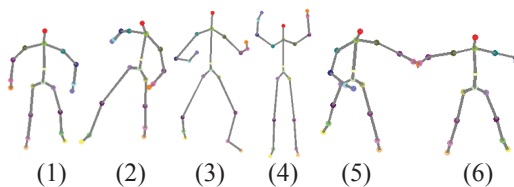


Figure 6: Six average poses of individual clusters: our latent variable encodes pose states and reduce the pose diversity in each cluster. Average poses (1), (4) and (6) show the hands changes while (1), (2) and (5) focus on the facing direction.

211

212 4.2. Learning of the model

213 Random forests [26] have been proved as an effect and efficient algorithm
214 for human pose estimation on depth images. This section introduces how
215 to learn the structure of random forest and the corresponding parameters of
216 appearance models. We learn the structure of random forest based on the
217 method in [8], but different from [8], we treat each sample as a two-label
218 structure.

219 **Overview of random forest.** Random forest $\Gamma = \{T_t\}$ is a collection
220 of randomized decision trees T_t . Each tree T_t is built on a randomly selected

221 subset of training samples and learns a mapping from a sampled point to
 222 parameter space Θ . For the classification task, the parameter space is the
 223 label set, indicating the body part, and for the regression task, it may be R^3 in
 224 our case. To learn the structure of tree T_t , the selected samples corresponding
 225 to tree T_t will be iteratively divided into two separated subsets by a binary
 226 splitting function ζ . The splitting function ζ could be simple comparison of
 227 feature values and its threshold is generated randomly. The best one of the
 228 splitting functions will be chosen by maximizing the information gain. We
 229 use $S = \{s_i\}$ to denote the set of the training samples and S_L, S_R for the two
 230 split subsets. As a result, the destination function can be written as

$$\zeta^* = \arg \min_{\zeta} g(\zeta), \quad (5)$$

231

$$g(\zeta) = H(S) - \sum_{i \in \{L, R\}} \frac{|S_i|}{|S|} H(S_i), \quad (6)$$

232 where $H(\cdot)$ is the entropy or the sum-of-squared-differences depending on
 233 the specific task. This splitting continues recursively until the stop criteria
 234 are met, e.g. the tree reaches the maximal depth or there are less than a
 235 minimum number of samples in set S .

236 **Learning tree structures.** We treat each pixel labeled by a body part
 237 on the depth image as a sample and use random forest for the multi-label
 238 classification task. If each sample subset is used to train each sub-model
 239 independently, the complexity of the final model will increase at least linearly

240 in number of states of the latent variable. To address this issue, we employ
 241 a shared-structure model to train the random forest. We see each sample
 242 (pixel) as a multi-tag object $s_i = (f_i, l_i, c_i)$, where f_i refers to features, l_i
 243 refers to the body part label and c_i refers to the latent state. To fit the
 244 multi-tag samples, we adjust the expression of entropy $H(S)$ to be

$$H(S) = \sum_{c \in C} H(S_c), \quad (7)$$

$$H(S_c) = - \sum_{l_i} p_{l_i, c} \log(p_{l_i, c}), \quad (8)$$

245 where $H(S_c)$ is the entropy from the sample subset under the same latent
 246 state c , and $p_{l_i, c}$ is the probability of the sample with the label l_i in the
 247 subset. We adopt the depth comparison features proposed in [6], then the
 248 splitting function ζ for sample s could be:

$$\zeta(s; k, \eta) = \begin{cases} 0, & \text{if } f_s(k) < \eta, \\ 1, & \text{otherwise.} \end{cases} \quad (9)$$

249 where $f_s(k)$ is the k th value in the depth comparison features and η is the
 250 random threshold.

251 **Parameters of appearance model.** At each leaf ι of a tree we learn
 252 a compact expression $p(x_i | \iota, c_k)$ of votes for the position x_i conditional on
 253 the value of latent variable c_k . Specifically, for each sample set with latent

254 variable c_k , a mean-shift algorithm with a Gaussian kernel is applied to clus-
 255 ter the relative votes which present the offsets from the sampled position
 256 to the body part. The largest M centers $\{\Delta_{\iota mc_k}\}$ are stored at leaf node ι
 257 with a confidence weight $w_{\iota mc_k}$ which is equal to the size of the cluster. As a
 258 result, the conditional distribution $p(x_i|\iota, c_k)$ can be expressed by using the
 259 Gaussian Parzen density estimators as:

$$p(x_i|c_k, \iota) \propto \sum_{m \in M} w_{\iota mc_k} \exp\left(-\frac{\|x_i - (\Delta_{\iota mc_k} + x_s)\|^2}{b^2}\right), \quad (10)$$

260 where x_s is the 3D location of sampled point s , b is the kernel bandwidth
 261 and we set an empirical value 0.05m in the experiments. While (10) models
 262 the probability for a voting element arriving at the leaf ι of a single tree, the
 263 probability over the forest is calculated by averaging over all trees,

$$\phi(x_i|c_k) \propto \frac{1}{|T|} \sum_{T_t \in T} p(x|c_k, \iota_t), \quad (11)$$

264 where ι_t is the corresponding leaf of tree T_t in the forest.

265 **Parameters of prior model.** Besides learning parameters of the ap-
 266 pearance model at each leaf ι , we also learn a compact expression $p(\Delta_{ij}|\iota, c_k)$
 267 of the relative position between joints i and j conditional on the value of
 268 latent variable c_k using the similar method as that in the learning of ap-
 269 pearance parameters. We use $\{\Delta_{ij, \iota mc_k}\}$ to denote the learned centers of the
 270 relative position between joints i and j by mean-shift algorithm and $w_{ij, \iota mc_k}$

271 to denote its weight. So, the relative position distribution between joints i
 272 and j conditional on the leaf ι and latent variable c_k can be expressed as

$$p(x_i, x_j | c_k, \iota) \propto \sum_{m \in M} w_{ij, \iota m c_k} \exp\left(-\frac{\|x_i - x_j - \Delta_{ij, m c_k}\|^2}{b_{ij}^2}\right), \quad (12)$$

273 where x_i and x_j are the estimated positions of joints i and j , and b_{ij} is the
 274 kernel bandwidth, which we set to the average limb length in the training
 275 data. As a result, the probability of the forest is calculated by averaging over
 276 all trees,

$$\phi(x_i, x_j | c_k) \propto \frac{1}{|T|} \sum_{T_t \in T} p(x_i, x_j | c_k, \iota). \quad (13)$$

277 Compared with the Gaussian prior model, our prior model builds its expres-
 278 sion using specific sampling points on each test image. This would enhance
 279 adaptability of a prior model.

280 5. Experiments and Discussion

281 5.1. Datasets

282 In this section, we evaluate our algorithm for human pose estimation on
 283 two depth datasets, the Stanford dataset [12] and our THU pose dataset.

284 **The Stanford dataset.** It consists of 28 action sequences of one person,
 285 which includes 7891 images in total with a resolution of 176×144 . The images
 286 were captured by using a ToF camera in a lab environment and joint positions
 287 are obtained by motion sensors. Among the images, 6000 are selected for
 288 training and the rest, less than 2000, are for testing.

289 **The THU dataset2.** To further evaluate our method, we collect a new
290 dataset for experiments. Our THU dataset2 contains 15000 depth images
291 captured by a Kinect camera, which consists of 5 persons performing general
292 actions (including upper/lower limbs movements, turning, jumping, etc.).
293 Some samples are shown in Fig. 7. We use motion detection method, such
294 as [27, 28], to get the foreground manually labeled landmarks as the ground
295 truth. Among the images, 10000 are randomly selected for training and the
rest are for testing.

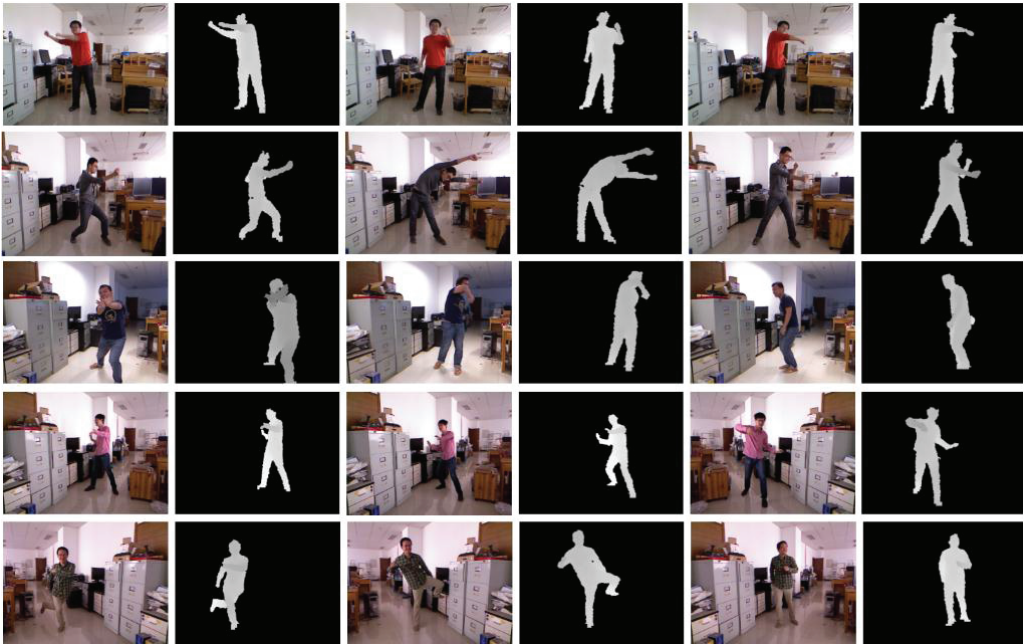


Figure 7: Samples from the THU dataset2: RGB and depth images.

296

297 *5.2. Preprocessing of the training data*

298 We assume the foreground is clear in our model. So to ensure this, some
299 preprocessing should be done before training. We perform a motion-based
300 method [29] to segment the foreground from background. Some segmentation
301 results in the Stanford dataset are shown in Fig. 8. Besides, the baseline
302 method [8] used in this paper needs to label the pixels for each part. It
303 involves a great deal of work. To facilitate this, we label each pixel as the
nearest body part.



Figure 8: Results of foreground segmentation of [25] in the Stanford dataset: pairs of original and foreground images.

304

305 *5.3. Performance evaluation*

306 To evaluate our algorithm, we compare our proposed method with some
307 state-of-the-art methods in [8, 30, 12, 11]. Two measures are used to demon-
308 strate the performance: the average error and the mean of average precision
309 (mAP). The average error for each joint evaluates the average difference be-
310 tween the estimated position and its ground truth under the Euclidean space
311 and the mAP presents the ratio of the most confident joint hypothesis within
312 the distance tolerance $\tau = 0.1\text{m}$, as with [8]. For the specific joint i , its mAP
313 can be calculated by

$$mAP_i = \frac{1}{M} \sum_{m=1}^M 1(|\hat{x}_i(m) - x_i(m)| < \tau), \quad (14)$$

314 where M is the number of testing samples, $\hat{x}_i(m)$ is the estimated position
 315 of joint i , $x_i(m)$ is the ground-truth and $1(\cdot)$ is an indicator function.

316 **Experiments on the Stanford Dataset.** Considering the sample size
 317 and pose variation in this dataset, we set K to 4 ($\|C\| = 4$), the centers of
 318 the four clusters are shown in Fig. 9, and we set $T = 0.2$ for the inference
 319 stage. The influence of the cluster number K and the threshold T will be
 320 discussed in Section 5.4.

321 On this dataset, we compare our method with some state-of-the-art meth-
 322 ods [8, 30, 12, 11]. The experimental results are shown in the second column
 323 of Table 1. We can observe that compared with the published results, our
 324 method obtains a better result, the mAP of 98.2%. Some of the estimated
 325 results are illustrated in Fig. 10. From Fig. 10, it can be found that our
 326 method can get good results for the most samples with a front-facing an-
 327 gle and with a small side-facing angle. We note that it fails under a large
 328 side-facing angle, the results are shown in the black box in Fig. 10. It is a
 329 challenging task to estimate human pose within a side-standing body. The
 330 first result in the black box shows that our method fails to estimate the part
 331 on the right body due to a large area occlusion. The second result in the
 332 black box shows that our method makes a symmetric error because it cannot
 333 recognize a back-facing body on this depth image. To overcome this issue,

334 methods in [31, 32] on sequence of images or some tracking methods in [33]
 335 may help. Additionally, we test the speed of our algorithm in processing
 336 one image on the Stanford dataset. With our non-optimized code, it runs
 337 the processing at about 36fps on our 4-cores computer. This would be fast
 enough for many visual interaction tasks.

Table 1: Comparison of mAP ($\tau = 0.1m$) with some state-of-the-art methods.

Method	On Stanford dataset	On THU dataset2
Ganapathi et al. [12]	0.898	–
Ye et al. [11]	0.950	–
Shotton et al. [6]	0.947	–
Girshick et al. [8]	0.957	0.89
He15 [25]	0.98	0.88
ours	0.982	0.971

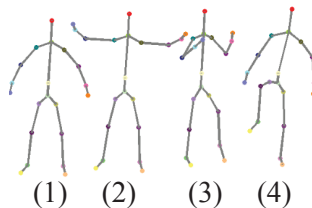


Figure 9: The centers of clusters on the Stanford dataset.

338

339 **Experiments on the THU Dataset2.** For this dataset, we set K to
 340 16 and the average poses are shown in Fig. 11. For each cluster, we use the
 341 method in [25] to train the random forest. The rest of the settings are the
 342 same as that on the Stanford dataset.

343 We compare our approach to a state-of-the-art method proposed by Gir-
 344 shick et al. [8] and the method in [25]. They both estimate the joint locations

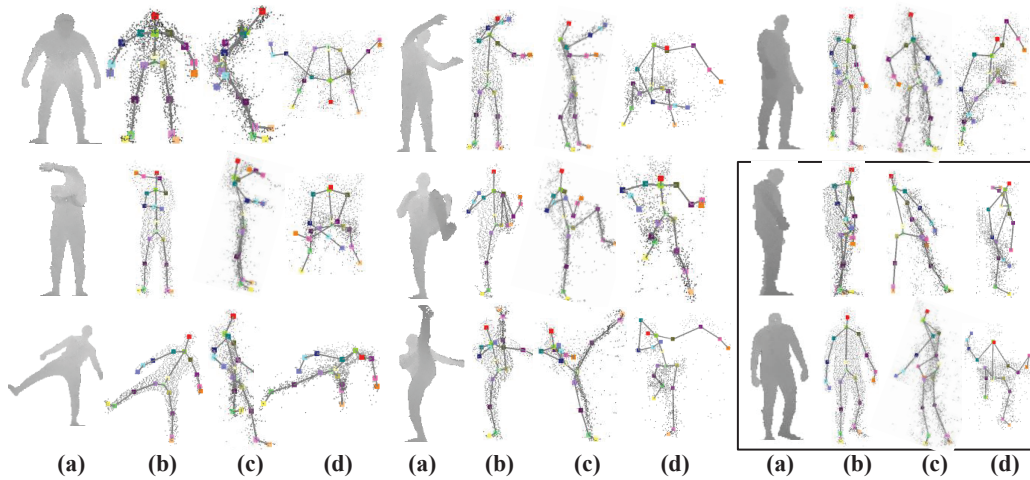


Figure 10: Nine estimated results on the Stanford dataset: (a) the original depth image; (b), (c) and (d) our results from the front view, the left-side view and the top view, respectively. Results in the box are those that our method fails.

345 by regression forest. Experimental results are shown in the third column of
 346 Table 1. The detailed comparison of the approach [8], denoted by ‘Girshick et
 347 al.’, and our LVPS, denoted by ‘ours’, is shown in Fig. 12. From the Fig. 12,
 348 we can find that our algorithm achieves better results than that of [8]. More
 349 specifically, our algorithm obtains 3.6cm in the average error and 97.1% in
 350 mAP. Besides, the superior results can be remarkably observed at limb ends,
 351 such as elbow, wrist and hand, which we think benefits from the use of la-
 352 tent models and the graphical models. Compared with the method [25], our
 353 method yields a better result. By the way, the method [25] can be seen as
 354 the case that $K = 1$ the proposed algorithm. Some samples are illustrated
 355 in Fig. 13.

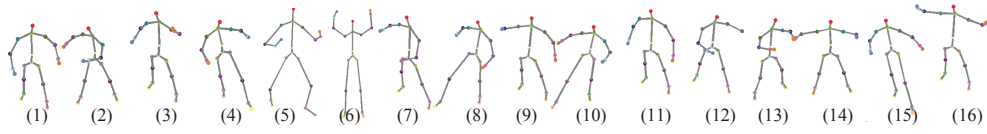


Figure 11: The centers of clusters on the THU dataset2.

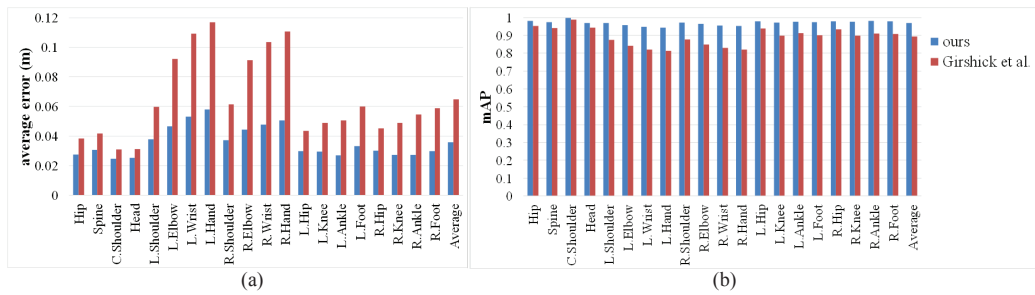


Figure 12: Performance on the THU dataset2: (a) average estimation error vs. body joint; (b) mAP vs. body joint.

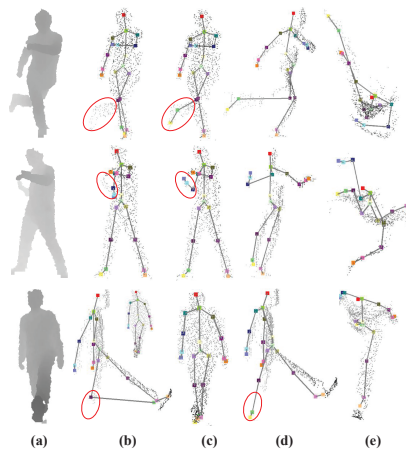


Figure 13: Three estimated sample images from the method in [8] and ours: (a) the original depth image; (b) results from the method [8]; (c), (d) and (e) our results from the front view, the left-side view and the top view, respectively.

356 *5.4. Discussion*

357 In this section we investigate the effects of three main factors that may
358 affect the pose estimation accuracy of our method. These factors are the
359 cluster number $K = \|C\|$, the construction of the inference model and the
360 threshold T .

361 **Cluster Number K .** We retrain our models with different cluster num-
362 bers K from 1 to 32 on both datasets. The results are shown in Fig. 14. On
363 the THU dataset2, when K is increased from 1 to 16, the value of mAP is
364 enhanced from about 0.88 to 0.97 and after that it drops. It illustrates that
365 the larger the cluster number K is, the better the models are learned, but if
366 K is too large, it causes over-fitting. On the Stanford dataset, splitting the
367 pose space does not boost the performance. We think the small diversity of
368 the pose on the Stanford dataset causes this. Nevertheless, when K is equal
369 to 1, the method can be seen as the method [25]. Compared with it, we can
370 observe the superiority of our method.

371 **Construction of Inference Model.** In the inference stage, we use a
372 linear strategy (3) to construct the detection model. We compare our strategy
373 in (3) with another usual strategy: using a fixed value of K^* ($K^*=1$ and
374 2) for inference, denoted by ' $K^* = 1$ ' and ' $K^* = 2$ '. $K^*=1$ means the most
375 plausible sub-model is used for inference while $K^*=2$ means that two sub-
376 models with the largest likelihoods are linearly combined for inference. The
377 results are shown in Table 2. We can observe that our proposal obtains the
378 best result among these combining methods, which indicates the effectiveness

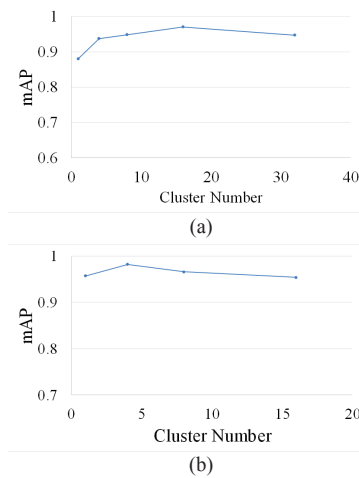


Figure 14: mAP vs. cluster number: (a) results on the THU dataset2, (b) results on the Stanford dataset.

379 of our method. Moreover, it can be observed that the results of ‘ $K^* = 1$ ’ and
 380 ‘ $K^* = 2$ ’ are very close. It implies that our latent variable is discriminative
 to cluster the pose.

Table 2: Performance of various combining strategies.

Method	mAP ($\tau = 0.1m$)
$K^* = 1$	0.956
$K^* = 2$	0.962
ours	0.97

381

382 **Threshold T .** The threshold T controls the number of sub-models used
 383 for inference. We investigate the pose estimation performance under different
 384 thresholds T and show the results in Fig. 15. Although it yields the best
 385 results at $T = 0.2$, it still maintains an mAP of higher than 0.9 for other
 386 values of T , which indicates the robustness of our model. Additionally, in
 387 order to further show how the threshold T works, we calculate the proportions

388 of cluster numbers used for inference in Fig. 16. It demonstrates that as the
 389 threshold T goes up, there are more clusters used for inference. Before T
 390 reaches 0.2, only the most probable model is used to estimate the human pose.
 391 After that, more and more models are involved in the inference. Considering
 392 both the results in Fig. 15 and Fig. 16, we find that merging the proper
 number of models can improve the performance.

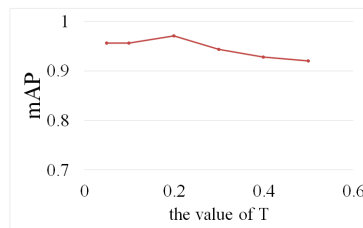


Figure 15: mAP vs. the threshold T .

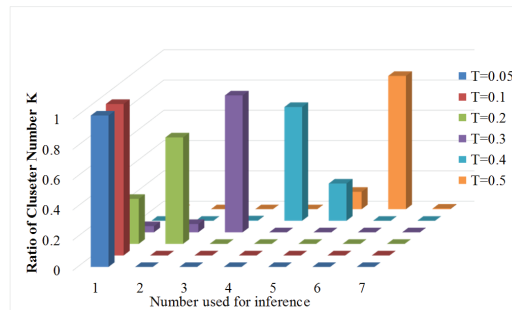


Figure 16: The proportions of cluster numbers used for inference under different thresholds T : different colors indicate the values of T .

393

394 **6. Conclusion and Future Work**

395 In this paper, we have proposed a novel approach to pose estimation on
396 depth images. In the approach, we have proposed the latent variable pictorial
397 structure (LVPS) to adapt the prior model and enhance the discrimination
398 of the appearance model by incorporating a latent variable. We have also
399 modified the silhouette features to encode the human pose, clustered the
400 pose space and established a new pose dataset to evaluate the performance
401 of the proposed method. Through these enhancements, our LVPS model can
402 learn better appearance and prior models. Experiments have verified that the
403 proposed method could achieve higher accuracy on the published datasets,
404 compared with other state-of-the-art methods. It would be interesting to
405 further our work by combining this method with object tracking.

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