Simple Pose: Rethinking and Improving a Bottom-up Approach for Multi-Person Pose Estimation

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Abstract

We rethink a well-known bottom-up approach for multi-person pose estimation and propose an improved one. The improved approach surpasses the baseline significantly thanks to (1) an intuitional yet more sensible representation, which we refer to as body parts to encode the connection information between keypoints, (2) an improved stacked hourglass network with attention mechanisms, (3) a novel focal L2 loss which is dedicated to “hard keypoint and keypoint association (body part) mining, and (4) a robust greedy keypoint assignment algorithm for grouping the detected keypoints into individual poses. Our approach not only works straightforwardly but also outperforms the baseline by about 15% in average precision and is comparable to the state of the art on the MS-COCO test-dev dataset. The code and pre-trained models are publicly available online.¹

Introduction

The problem of multi-person pose estimation aims at recognizing and localizing the anatomical keypoints (or body joints) of all persons in a given image. Considerable progress has been made in this field, benefiting from the development of more powerful convolutional neural networks (CNNs), such as ResNet (He et al. 2016) and DenseNet (Huang et al. 2017), and more representative benchmarks, such as MS-COCO (Lin et al. 2014).

Existing approaches tackling this problem can be divided into two categories: top-down and bottom-up. The top-down approaches, e.g. (Gao et al. 2017; Papandreou et al. 2019; Sun et al. 2019), usually employ a state-of-the-art (SOTA) detector, such as SSD (Liu et al. 2016), to capture all the persons from the image first. Then, the cropped persons are resized and fed into the SOTA pose estimator designed for a single person, e.g. (Wei et al. 2016; Newell, Yang, and Deng 2016; Chen et al. 2018). In contrast, the bottom-up approaches, e.g. (Papandreou et al. 2017; Kreiss, Bertoni, and Alahi 2019), directly infer the keypoints and the connection information between keypoints of all persons in the image without a human detector. Afterwards, the keypoints are grouped to form multiple human poses based on the inferred connection information.

Top-down approaches, e.g. (Papandreou et al. 2017; Sun et al. 2019; Li et al. 2019), usually have complicated structures and low performance-cost ratios. Compared with them, the bottom-up approaches, e.g. (Newell, Huang, and Deng 2017; Kreiss, Bertoni, and Alahi 2019), can be more efficient in inference and independent of human detectors. However, they have to group the keypoints correctly. And the keypoint grouping (pairing or association in other words) can be a big challenge, resulting in another bottleneck for real-time usage (Pishchulin et al. 2016; Iqbal and Gall 2016). CMU-Pose (Cao et al. 2017), PersonLab (Papandreou et al. 2018) and PifPaf (Kreiss, Bertoni, and Alahi 2019) use the greedy parsing algorithm to group detected keypoints into individual poses and break through the bottleneck to some extent.

It is worth mentioning that the approach proposed by (Cao et al. 2017) (referred to as CMU-Pose here for convenience) is the first bottom-up approach to perform the task of multi-person pose estimation in the wild with high accuracy, and almost in real time. Our approach is mainly inspired by this work but is more intuitive yet more powerful. Hence, CMU-Pose is selected to be the baseline approach in this work.

The main contributions of this paper are summarized as follows: (1) we rethink the encoding of joint association, which is named as Part Affinity Fields (PAFs) (Cao et al. 2017), and propose a simplified yet more reasonable one, which we call body parts, (2) we present an improved stacked hourglass network with attention mechanisms to generate high-res and high-quality heatmaps, (3) we design a novel loss to help the network learn “hard samples, and (4) we develop the greedy keypoint assignment algorithm.

Related Work and Rethinking

Single Person Pose Estimation

Classical approaches tackling the problem of person pose estimation are mainly based on the pictorial structures (Fischler and Elschlager 1973; Andriluka, Roth, and Schiele 2009) or the graphical models (Chen and Yuille 2014). They usually formulate this problem as a tree-structured or graphical model problem and detect the keypoints based on hand-crafted features (Chen et al. 2018). Recently, the SOTA approaches leverage advanced CNNs and more abundant datasets, making enormous progress in pose estimation.

¹https://github.com/jialee93/Improved-Body-Parts
Here, we mainly discuss the CNN based approaches.

DeepPose (Toshev and Szegedy 2014) employs CNNs to solve this problem for the first time, by regressing the Cartesian coordinates of the joints (or keypoints) directly. By contrast, the work (Tompson et al. 2014) presents CNNs to firstly predict the Gaussian response heatmaps of keypoints, and subsequently it obtains the keypoint positions via finding the local maximums in the heatmaps. Some up to date work, e.g. (Papandreou et al. 2018; Kreiss, Bertoni, and Alahi 2019), decomposes the problem of keypoint localization into two subproblems at each pixel location: (1) binary classification (0 or 1), and (2) regression of offset vector to the nearest keypoint.

Multi-Person Pose Estimation

Top-down approaches. Most of the SOTA results have been achieved by the top-down approaches, such as CPN (Chen et al. 2018) and HRNet (Sun et al. 2019). Benefitting from the existing well-trained person detectors, the SOTA top-down approaches bypass the difficult subproblem of human body detection and turn the detection challenges into their advantages. However, they depend on the human detector heavily and they perform the task in two separate steps. The inference time will significantly increase if many people appear together.

Bottom-up approaches. The bottom-up approaches, e.g. (Cao et al. 2017; Newell, Huang, and Deng 2017; Kreiss, Bertoni, and Alahi 2019), are more efficient in keypoint inference and do not rely on the human detector. However, they tend to be less accurate. One main reason is that too large and too small persons in the image are difficult to detect at the same time (pose variation and feature map down-sampling make things even worse). Another main reason lies in the fact that the offset of only a few pixels away from the annotated keypoint location can lead to a big drop in the evaluation metrics (Wang et al. 2018) on the MS-COCO benchmark (Lin et al. 2014). On the contrary, the top-down approaches are immune to these challenges.

Some Rethinking

The topic of scale invariance in pose estimation is of great importance. Image pyramid (or multi-scale search in other words) technique is usually employed during testing to cover the human poses of different scales as much as possible (Cao et al. 2017; Newell, Huang, and Deng 2017; Papandreou et al. 2018), while the network is supervised at relatively a smaller scale range during training. Besides, some related work, e.g. (Newell, Yang, and Deng 2016; Chu et al. 2017; Ke et al. 2018), has designed special model structures to enhance the model invariance across scales.

High-res and high-quality feature maps (include the output heatmaps) are critical for accurate keypoint localization. Offset regression in PersonLab (Papandreou et al. 2018) and CornerNet (Law and Deng 2018), integral pose regression (Sun et al. 2018), and retaining (Sun et al. 2019) or even magnifying (Wang et al. 2018) the resolution of feature maps through the network are all good tries to relieve the precision loss (can not be avoided) caused by image or feature map resizing and small input or feature map size.

The encoding of connection (or association) information between keypoints is paid a lot of attention in some prior work, e.g. (Cao et al. 2017; Newell, Huang, and Deng 2017; Kreiss, Bertoni, and Alahi 2019; Papandreou et al. 2018). New representations bring about new ways of addressing problems. In this work, we only review the encoding of joint association named Part Affinity Fields (Cao et al. 2017) due to the limited space.

Another topic worthy of mention is the problem of imbalanced data: “positive samples vs “negative samples (between classes) and “easy samples vs “hard samples (within classes). A (Gaussian response) heatmap has most of its area equal to zero (background) and only a small portion of it corresponds to the Gaussian distribution (foreground). Thus, the spread of the Gaussian peaks should be controlled properly to balance the foreground and background. On the other hand, too many easy samples (such as Gaussian peaks of facial keypoints and easy background pixels) can prevent the network from learning the “hard” samples (such as Gaussian peaks corresponding to occluded keypoints or body parts) well. These two types of data imbalance problems are critical and should be addressed properly.

Our Approach

We perform the task in three steps: (1) we predict the keypoint heatmaps and the body part heatmaps of all persons in a given image, (2) we get the candidate keypoints and body parts by performing Non-Maximum Suppression (NMS) on
the inferred heatmaps, and (3) we perform the keypoint assignment algorithm and collect all individual poses.

**Definition of Heatmaps**

Considering the vague concept of keypoints and body parts, and human annotation variances (jitters), our network is supervised to regress the Gaussian responses (0~1 values) around the keypoint or body part area before obtaining the final localization, introducing the smooth mapping regularization and forcing the network to learn more features nearby.

The heatmaps here include keypoint heatmaps and body part heatmaps. Each pixel value in the keypoint heatmaps encodes the confidence that a nearest keypoint of a particular type occurs. We generate the ground truth keypoint heatmaps by putting unnormalized Gaussian distributions with a standard deviation $\sigma_k$ at all annotated keypoint positions. For example, the generated Gaussian peak of the right shoulder (RS) keypoint depicted in the Left of Figure 2.

The body part here refers to the body area which lies between the two adjacent keypoints (for instance, the forearm area in the Left of Figure 2). A set of body parts are used to encode the connection information between keypoints and extract the visual patterns of human skeleton (see the Middle of Figure 2). Since body part segmentations are not available and person masks only cover visible human body areas, we use an elliptical area to approximately represent the body part. We generate the ground truth body part heatmaps by putting unnormalized elliptical Gaussian distributions with a standard deviation $\sigma_p$ in all body part areas.

By the way, we map the pixel $p$ at the location $(x, y)$ in the $j$-th ground truth heatmap to its original floating point location $\tilde{p}(\tilde{x}, \tilde{y}) = \tilde{p}(x \cdot R + R/2 - 0.5, y \cdot R + R/2 - 0.5)$ in the input image, in which $R$ is the output stride, before generating the precise ground truth Gaussian peaks.

The PAFs proposed by (Cao et al. 2017) is a 2D vector field for each pixel in the limb area, which encodes the location and direction information of a limb. All the pixels within the approximate limb area (may include outliers of the limb) have the same ground truth value, which brings about vagueness or even conflicts to the information representation. The body part representation is more sensible and composite. Pixels near to the major axes of the body parts have higher confidence and vice versa. And we only need the half dimensions of PAFs to encode the keypoint connection information, reducing the demand for model capacity.

During training, a single loss supervises the network to infer two kinds of Gaussian peaks, which have similar visual patterns and share the same formulation.

The standard deviations, $\sigma_k$ and $\sigma_p$, control the spread of the Gaussian peaks and they should be set properly to balance the foreground pixels and background pixels. The hyper-parameters $r_0$ and $d_0$ (see their meanings in Figure 2) determine the boundaries of the ground truth Gaussian peaks, truncating the unnormalized Gaussian distribution at a fixed value $\theta_r$. It plays a role in our loss function.

**Network Structure**

Large receptive fields in CNN are critical for learning long range spatial relationships and can bring about accuracy improvement (Wei et al. 2016). On the other hand, the detailed information (in smaller receptive fields) is needed for fine-grained localization. To consolidate the global and local features, hourglass networks (Newell, Yang, and Deng 2016; Newell, Huang, and Deng 2017) have been designed to capture the different spatial extent information of each keypoint and association between keypoints by repeated bottom-up and top-down inference. In this work, we select the hourglass network, designed for multi-person pose estimation in Associative Embedding (AE) (Newell, Huang, and Deng 2017), as the base model and present an improved one. The improved variant, which we call Identity Mapping Hourglass Network (IMHN), significantly outperforms the hourglass network in AE (Newell, Huang, and Deng 2017).

The proposed IMHN, whose structure is depicted in Figure 3, is fully convolutional. It takes an image of any shape as the input and outputs multi-scale keypoint and body part heatmaps of all persons (if any) in the scene simultaneously. Before fed into the stacked hourglass modules, the original input is down-sampled twofold by some convolutional layers and max-pooling layers.

A first order hourglass module is designed as shown in Figure 4. The down-sampling path reduces the spatial extent of the input feature map by half once and increases the number of the feature map channels $C$ by $N$ ($C = 256$ and $N = 128$ in all experiments unless mentioned otherwise). After replacing the dashed box in Figure 4 with another first order module, we get a second order module. A fourth order module can be made by repeating this operation and it is the default hourglass module to build the IMHN. Therefore, the feature map in the deepest path
Loss Functions

The L2 loss is frequently used to measure the distance between the predicted heatmaps and the target heatmaps, e.g. (Wei et al. 2016; Cao et al. 2017; Ke et al. 2018). To handle the “hard” keypoints, the work (Chen et al. 2018) proposes the L2 loss with online “hard” keypoint mining. Here, we present a novel loss, which we refer to as focal L2 loss, under the unified definition of keypoint and body part heatmaps, to deal with the two types of sample imbalance problems as introduced in Section Some Rethinking.

At each stage of the stacked IMHN, $K$ keypoint heatmaps and $P$ body part heatmaps are inferred at 5 different scales. A pixel value in the inferred heatmaps represents the confidence of being a certain category of keypoint or body part. Assuming the predicted score maps (or heatmaps$^2$) of size $w_i \times h_i$ at stage $t$ are $S^t = (S^{t_1}_1, S^{t_2}_2, \ldots, S^{t_K}_{K+P})$, $t \in \{1, 2, \ldots, T\}$, where $T$ is the total number of stacked hourglass modules. Supposing the ground truth heatmaps of the same size are $S^* = (S^*_1, S^*_2, \ldots, S^*_{K+P})$ and the Gaussian peak generation function is $G$. Let $S^*_j(p)$ denote the ground truth score at the pixel location $p(x, y) \in \mathbb{R}^{w_i \times h_i}$ in the $j$-th heatmap, we compute $S^t_j(p)$ as:

$$S^t_j(p) = \begin{cases} G(x, y; R, \sigma_k, r_0), & 1 \leq j \leq K \\ G(x, y; R, \sigma_p, t_0), & K < j \leq K + P. \end{cases}$$

We define $Sd^t_j(p)$:

$$Sd^t_j(p) = \begin{cases} S^t_j(p) - \alpha, & S^t_j(p) > \text{thre} \\ 1 - S^t_j(p) - \beta, & \text{else} \end{cases}$$

where $\text{thre}$ (mentioned in Section Definition of Heatmaps) is the threshold to distinguish between the foreground heatmap pixels and the background heatmap pixels, and $\alpha$, $\beta$ are compensation factors to reduce the punishment of easy samples (both easy foreground pixels and easy background pixels) so that we can make full use of the training data.

The focal L2 loss ($\mathcal{FL}_i$) between the predicted heatmaps and target heatmaps of size $w_i \times h_i$ at stage $t$ is computed as follow:

$$\mathcal{FL}_i = \frac{1}{K+P} \sum_{j=1}^{K+P} \sum_{p \in \mathbb{R}^{w_i \times h_i}} [\eta \cdot \mathbb{I}(j \leq K) + 1] \cdot W(p) \cdot \frac{\|S^*_j(p) - S^t_j(p)\|^2_2}{\|1 - Sd^t_j(p)\|^2_2}.$$
The candidate keypoints are assigned provided that the candidate body parts are assigned into corresponding human skeletons. We perform NMS (3 × 3 window) on the predicted heatmaps to find the candidate keypoints. Then, we obtain the candidate body parts that lie between the candidate adjacent keypoints, and calculate their scores that represent the confidence of being body parts, by sampling a set of Gaussian responses within the body part areas. After that, the candidate body parts of the same type are sorted in descending order according to the weighted scores of body parts and connected keypoints. Consequently, K sets of keypoints \( J = \{J_s_1, J_s_2, \ldots, J_s_K\} \) and \( P \) sets of body parts \( L = \{L_s_1, L_s_2, \ldots, L_s_P\} \) are obtained.

Each element \( l_{i,j} \in L_s_i \) is a body part instance with type ID \( i \), connected keypoints and weighted score \( S_{h,j} \). Here, one candidate keypoint cannot be shared by two or more body parts of the same type, i.e., \( \forall j, k \) and \( m \), \( \{l_{i,j}, l_{i,k}\} \subseteq L_s_i \) and \( j \neq k \), we ensure that \( l_{i,j} \cap l_{i,k} \cap J_s_m = \emptyset \), where \( J_s_m \in J \). This rule works in analogy to the NMS for candidate bounding boxes in object detection task.

Supposing the set of assembled human poses is \( H = \{h_1, h_2, h_3, \ldots\} \), in which \( h_n \in H \) represents a single person pose. Then, \( h_n \) has a \( 1 \sim P \) assigned body parts, corresponding type IDs and total score \( S_{h_n} \). Our goal is to select the proper candidate body parts and find the best grouping strategy between the body parts in \( L_s_1, L_s_2, \ldots, L_s_P \), such that \( \sum_{h_n \in H} S_{h_n} \) reaches to its global maximum. Instead of solving the problem of global graph matching globally, CMU-Pose (Cao et al. 2017) proposes a greedy algorithm upon a minimum spanning tree (MST) of human skeleton to match the adjacent tree nodes independently at only a fraction of the original computational cost.

We follow the greedy strategy in CMU-Pose and assemble the human skeletons (see the Middle of Figure 2) by matching adjacent body parts independently. Our keypoint assignment algorithm is based on several simple connection rules. As redundant body parts are introduced in our human skeleton, the assembled body parts having lower scores are removed by the body parts having higher scores, which share the same connected keypoint(s).
Table 1: Results on the MS-COCO 2017 validation set.

| ID | Method                        | Input Stride | FL   | AP  | ID | Method                        | Input Stride | FL  | AP  |
|----|-------------------------------|--------------|------|-----|----|-------------------------------|--------------|------|-----|
| 1  | CMU-Pose (6-stage CMU-Net)    | 368 times 8  | N    | 56.0| 12 | 3-stage IMHN, w/ MST          | 384 times 4  | Y   | 65.1|
| 2  | AE (4-stage Hourglass, + val data) | 512 times 4  | N    | 59.7| 13 | 4-stage IMHN                  | 384 times 4  | N   | 64.5|
| 3  | Top-down* (8-stage Hourglass) | 256 times 4  | N    | 66.9| 14 | 4-stage IMHN                  | 384 times 4  | Y   | 67.3|
| 4  | Ours (3-stage CMU-Net)        | 368 times 8  | N    | 56.5| 15 | 4-stage IMHN, + val data      | 384 times 4  | Y   | 72.3|
| 5  | Ours (3-stage CMU-Net)        | 368 times 8  | Y    | 60.7| 16 | 4-stage IMHN plus, + val data | 512 times 4  | Y   | 69.1|
| 6  | Ours (4-stage Hourglass)      | 512 times 4  | N    | 60.0| 17 | 4-stage IMHN plus              | 512 times 4  | Y   | 74.1|
| 7  | 3-stage IMHN                  | 384 times 4  | N    | 61.5| 18 | 4-stage IMHN, one scale       | 768 times 4  | Y   | 63.4|
| 8  | 3-stage IMHN                  | 384 times 4  | Y    | 65.8| 19 | 4-stage IMHN plus, one scale  | 768 times 4  | Y   | 65.9|
| 9  | 3-stage IMHN, w/o spatial attention | 384 times 4  | Y    | 64.6| 20 | PersonLab*, one scale         | 801 times 8  | Y   | 65.7|
| 10 | 3-stage IMHN, w/o channel attention | 384 times 4  | Y    | 65.4| 21 | PersonLab*, one scale         | 801 times 8  | Y   | 61.2|
| 11 | 3-stage IMHN, FL only for keypoint | 384 times 4  | Y    | 64.2| 22 | PersonLab*, one scale         | 1401 times 8 | Y   | 66.5|

Table 2: Results on the MS-COCO 2017 test-dev set.

| Method                        | Backbone            | Pretrain | Train Input | Test Input | Refine | AP  | AP* | AR  | AR* |
|-------------------------------|---------------------|----------|-------------|------------|--------|-----|-----|-----|-----|
| Bottom-up: multi-person keypoint detection and grouping |                      |          |             |            |        |     |     |     |     |
| CMU-Pose (baseline) (Cao et al. 2017) | 6-stage CMU-Net   | N        | 368 times 368 | ~ 368^2 | N  | 52.9 | 50.9 | 57.2 | 57.0 | 79.2 |
| CMU-Pose* (Cao et al. 2017)    | 6-stage CMU-Net   | N        | 368 times 368 | ~ 368^2 | Y  | 61.8 | 57.1 | 68.2 | 66.5 | 87.2 |
| AE* (Newell, Huang, and Deng 2017) | 4-stage Hourglass | N        | 512 times 512 | ~ 512^2 | Y  | 65.5 | 60.6 | 72.6 | 70.2 | 89.5 |
| PersonLab* (Papandreou et al. 2018) | ResNet-101      | Y        | 801 times 801 | ~ 1401^2 | N  | 67.8 | 63.0 | 74.8 | 74.5 | 92.2 |
| PifPaf (Kreiss, Bertoni, and Alahi 2019) | ResNet-101      | Y        | 401 times 401 | ~ 641^2  | N  | 64.9 | 60.6 | 71.2 | 70.3 | 90.2 |
| PifPaf* (Kreiss, Bertoni, and Alahi 2019) | ResNet-152      | Y        | 401 times 401 | ~ 641^2  | N  | 66.7 | 62.4 | 72.9 | 72.2 | 90.9 |
| Ours-1, w/ FL                  | 3-stage CMU-Net   | N        | 368 times 368 | ~ 368^2 | N  | 59.3 | 56.2 | 63.8 | 63.5 | 84.6 |
| Ours-2, w/ FL                  | 3-stage IMHN     | N        | 384 times 384 | ~ 384^2 | N  | 65.2 | 63.7 | 68.5 | 69.8 | 87.7 |
| Ours-3, w/ FL                  | 4-stage IMHN     | N        | 384 times 384 | ~ 384^2 | N  | 66.2 | 66.4 | 71.6 | 71.2 | 88.6 |
| Ours-4 (final), w/ FL          | 4-stage IMHN plus | N        | 512 times 512 | ~ 512^2 | N  | 68.1 | 66.8 | 70.5 | 72.1 | 88.2 |
| Ours-5, w/ FL                  | 4-stage IMHN plus | N        | 512 times 512 | ~ 512^2 | N  | 67.6 | 64.5 | 72.6 | 71.3 | 87.6 |

Top-down: human detection and single-person keypoint detection

| Method                        | Backbone | Pretrain | Train Input | Test Input | Refine | AP  | AP* | AR  | AR* |
|-------------------------------|----------|----------|-------------|------------|--------|-----|-----|-----|-----|
| G-RMI* (Papandreou et al. 2017) | ResNet-101 | Y        | 353 times 257 | 353 times 257 | –   | 64.9 | 62.3 | 70.0 | 69.7 | 88.7 |
| CPN* (Chen et al. 2018)       | ResNet-Inception | Y        | 384 times 288 | 384 times 288 | –   | 72.1 | 68.7 | 77.2 | 78.5 | 95.1 |
| HRNet-W48* (Sun et al. 2019)  | HRNet-W48  | Y        | 384 times 288 | 384 times 288 | –   | 75.5 | 71.9 | 81.5 | 80.5 | 95.7 |

Implementation Details

Dataset and evaluation metrics. Our models are trained and evaluated on the MS-COCO dataset (Lin et al. 2014), which consists of the training set (includes around 60K images), the test-dev set (includes around 20K images) and the validation set (includes 5K images). The MS-COCO evaluation metrics, OKS-based\(^3\) average precision (AP) and average recall (AR), are used to evaluate the results.

Training details. The training images with random transformations are cropped and resized to the fixed spatial extent of 384 × 384. And the generated ground truth heatmaps have the size of 96 × 96. We implement both the 3-stage IMHN and 4-stage IMHN using Pytorch. To train the networks, we use the SGD optimizer with the learning rate of 1e-4 (multiplied by 0.2 for every 15 epochs), the momentum of 0.9, the batch size of 32 and the weight decay of 5e-4. We train the IMHNs with L2 loss first and then continue to train them with the focal L2 loss (FL) until the performance refuses to improve. The last but not least, our networks are in mixed precision to reduce the memory consumption and speed up the experiments. The implemented 4-stage IMHN (see Ours-3 in Table 2) can be trained at the speed of 33 FPS with 4 RTX 2080TI GPUs, and we train it for about 3 days.

Testing details. We first resize and pad the input image so that it fits the network. Then, the input image is inferred at multiple scales (for example, \(\times 0.5, \times 1, \times 1.5, \times 2\)) with flip augmentation. Next, the inferred heatmaps are averaged across scales (i.e., multi-scale search). After collecting the poses from the heatmaps via the keypoint assignment algorithm, we sort them in descending order according to their scores in heatmaps. To compare equally, we have run the released models of CMU-Pose (Cao et al. 2017), AE (Newell, Huang, and Deng 2017) and PifPaf (Kreiss, Bertoni, and Alahi 2019), obtaining their results without the refinement for detected person poses.

Results and Analyses

Results on the MS-COCO dataset. We have trained all our networks from scratch only with the MS-COCO data. The hyper-parameters of our approach are tuned according to the performance on the validation set. We compare our approach with the state of the art in Tables 1 and 2. The backbone of 4-stage IMHN has nearly the same number of convolutional layers as ResNet-101 (He et al. 2016). Thus, we specially compare our networks with those based on ResNet-101 backbone in PersonLab (Papandreou et al. 2018) and PifPaf (Kreiss, Bertoni, and Alahi 2019) impar-

\(^3\)OKS (Object Keypoint Similarity) defines the similarity between different human poses. Only the top 20 scoring poses are considered during evaluation.
tially. The entries marked with "*" in Tables 1 and 2 are results reported in their original papers.

The notations in our tables are explained as follows: "Input" denotes the long edge of the test image, "Stride" is the ratio of the input image size to the output feature map size, "MST" is short for minimum spanning tree of human skeleton without redundant body parts, "CMU-Net" represents the cascaded CNN used in CMU-Pose (Cao et al. 2017), "Hourglass" denotes the hourglass network, "Refine" indicates whether or not the result is additionally refined by a single person pose estimator. The entry named "Top-down" in Table 1 is a top-down approach cited from CPN (Chen et al. 2018). It employs a SOTA human detector and an 8-stage hourglass network for single person pose estimation.

**Inference speed.** The speed of our system is tested on the MS-COCO test-dev set (Lin et al. 2014).

- Inference speed of our 4-stage IMHN with $512 \times 512$ input on one 2080TI GPU: 38.5 FPS (100% GPU-Util).
- Processing speed of the keypoint assignment algorithm that is implemented in pure Python and a single process on CPU: 5.2 FPS (has not been well accelerated).

**Ablation studies.** Some examples of the qualitative comparison between our approach and CMU-Pose is illustrated in Figure 1. The detailed ablation experiments of our approach, numbered for clarity, are shown in Table 1. Experiments 1 and 4 use different encodings of keypoint association (PAFs and body parts respectively). They reveal that the encoding of body parts is better. According to experiments 2 and 6, our approach equipped with the same network and L2 loss is comparable to AE which utilizes validation data. The focal L2 loss can bring about a 3~4% AP improvement over the L2 loss (see experiment 5 vs 4, experiment 8 vs 7 and experiment 14 vs 13) under the same definition of heatmaps. Experiments 8 and 11 demonstrate doing "hard" keypoint and body part (keypoint association) mining meanwhile is better. The 3-stage IMHN can lead to around a 5% AP improvement compared with the 3-stage CMU-Net (see experiment 7 vs 4 and experiment 8 vs 5), and the 4-stage IMHN outperforms the 4-stage hourglass network (see experiment 13 vs 6) by a big margin, indicating IMHNS’ advantages.

The introduced “spatial attention” and “channel attention” mechanisms contribute 1.2% and 0.4% AP increase respectively to the 3-stage IMHN (see experiments 8, 9 and 10). The redundant connections in human skeleton bring about a 0.7% AP improvement according to the comparative experiments of 8 and 12. Bigger input size or feature map size and more stacks can improve the accuracy consistently according to the results in Table 1. Thus, we continue to train the model in experiment 14 with $512 \times 512$ input and more data augmentation, and obtain the model named “4-stage IMHN plus” in experiment 16. Further, the 4-stage IMHN is able to fit the MS-COCO train-val data at 74.1% AP (see experiment 17), indicating the big promotion space of our system.

**Comparisons with the state of the art.** All the models compared in the tables are evaluated without model ensemble. Some SOTA top-down approaches may outperform all the SOTA bottom-up approaches including ours, but they all depend on advanced human detectors and use very powerful networks for single person pose estimation. According to the results, it is safe to conclude that our approach outperforms the baseline by a big margin (experiment 5 vs 1 in Table 1 and Ours-1 vs the baseline in Table 2) and even surpasses the TOP-DOWN approaches with equal level backbones (please refer to “Top-down” in Table 1 and “G-RMI” in Table 2).

Our approach has achieved comparable or even better results on both single scales (see the experiments with IDs 18~22 in Table 1) and multiple scales (see the results in Table 2), compared with the latest SOTA bottom-up approaches, PifPaf and PersonLab, under fair conditions. However, PersonLab benefits greatly from the big input size (see experiment 22 vs 21 in Table 1). It can be seen that PifPaf and Personlab are superior to our approach in keypoint average recall (AR). But our approach is superior to them when it comes to the robustness to person scales. Both PersonLab and PifPaf drop over 10% from AP$^L$ metric to AB$^M$ metric (see the entries in Table 2), while our approach (Ours-4) performs well regardless of the person scales, be they middle scales (M) and large scales (L). What is more, most of other work benefits greatly from the networks pre-trained for the ImageNet classification task (Russakovsky et al. 2015), while we train networks from scratch (see experiment 17).

Referring to the results on the MS-COCO test-dev set in Table 2, the proposed techniques except for the designed IMHN bring about a 6.4% AP improvement (Ours-1 vs baseline). The 3-stage IMHN can lead to a 5.9% AP increase compared with the 3-stage CMU-Net (Ours-2 vs Ours-1). The 4-stage IMHN can further contribute a 1% AP improvement over the 3-stage IMHN. And our final model, “4-stage IMHN plus” with bigger input (means bigger output feature maps), brings about a 15.1% AP improvement in total over the baseline. Incidentally, our approach significantly outperforms CMU-Pose and AE, though they have additionally refined the results using a single person pose estimator.

**Conclusions**

In this paper, we rethink and develop a bottom-up approach for multi-person pose estimation. We provide some insights into valuable design choices: (1) doing hard sample mining of keypoint and keypoint association meanwhile, (2) using a powerful network to generate high-res and high-quality heatmaps, and (3) introducing the scale invariance across person scales, which are more critical to improve the performance. The experimental results have demonstrated the significant improvement achieved by our approach over the baseline (+15.1% AP on the MS-COCO test-dev dataset). To the best of our knowledge, our approach, which is straightforward and easy to follow, is the first bottom-up approach to provide both the source code and pre-trained models with over 67% AP on the MS-COCO test-dev dataset.

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