HAA4D: Few-Shot Human Atomic Action Recognition via 3D Spatio-Temporal Skeletal Alignment

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Abstract

Human actions involve complex pose variations and their 2D projections can be highly ambiguous. Thus 3D spatio-temporal or 4D (i.e., 3D+T) human skeletons, which are photometric and viewpoint invariant, are an excellent alternative to 2D+T skeletons/pixels to improve action recognition accuracy. This paper proposes a new 4D dataset HAA4D which consists of more than 3,300 RGB videos in 300 human atomic action classes. HAA4D is clean, diverse, class-balanced where each class is viewpoint-balanced with the use of 4D skeletons, in which as few as one 4D skeleton per class is sufficient for training a deep recognition model. Further, the choice of atomic actions makes annotation even easier, because each video clip lasts for only a few seconds. All training and testing 3D skeletons in HAA4D are globally aligned, using a deep alignment model to the same global space, making each skeleton face the negative z-direction. Such alignment makes matching skeletons more stable by reducing intra-class variations and thus with fewer training samples per class needed for action recognition. Given the high diversity and skeletal alignment in HAA4D, we construct the first baseline few-shot 4D human atomic action recognition network without bells and whistles, which produces comparable or higher performance than relevant state-of-the-art techniques relying on embedded space encoding without explicit skeletal alignment, using the same small number of training samples of unseen classes.

1. Introduction

The goal of human action recognition in computer vision is to classify the person’s action performed in a given video. Various approaches have been studied, such as RGB-based approaches [3, 6] which utilize color images and recently use CNN-based or LSTM-based methods for classification, or 2D+T approaches [13] that encode 2D skeletons into some latent space for traditional supervised learning. The use of 3D skeletons provide outstanding results compared to other methods, as 3D skeletons are photometric, geometric and viewpoint invariant, so that as few as one single 4D skeleton is sufficient to learn the pertinent action class (Figure 1). These invariances enable better temporal correlation by tracking the position of each 3D joint over time where each bone length is largely preserved throughout the action, in contrast to their projected 2D counterparts on images which may be distorted. Such invariances also make it easier for a computer algorithm to discern even fine differences in human poses.

Given the above advantages of 3D skeletons, previous human action dataset contributions [16, 25, 30, 31, 32] have provided valuable skeletal information for action recognition. However, these datasets suffer inherent disadvantages, such as lacking poses diversity or relatively low ground-truth accuracy. HAA4D is dedicated in complementing current human skeleton action datasets by providing a more comprehensive range of atomic actions in real-life scenarios, which have demonstrated excellent performance in recognizing composite and complex actions [9], while are easier to accurately annotate as each atomic action lasts for no more than 1–3 seconds. The samples in our dataset are collected from videos in-the-wild; therefore, we employ a deep
alignment model to predict the camera pose in each video frame, followed by aligning all 3D+T skeletons in an uniform camera coordinate system (Figure 1) such that all faces are facing in the negative $z$-direction. As human actions in the same class often share similar trajectories in their skeletal movements, such global alignment which transforms all train/test examples to the same coordinate system can significantly improve matching the pertinent action sequences using explicit geometry features.

With the emergence of deep learning and its superior performance in image classification tasks, related works mainly focus on supervised learning techniques that directly feed 4D human skeleton data into the network. Their networks are expected to generalize human actions and learn the underlying relationships in the embedded space. However, it is very difficult for deep learning to obtain satisfactory results with excessive learnable parameters when only a few examples are presented, given that 4D human skeletons are expensive to acquire. Furthermore, if the network is to recognize as many different human actions as possible, the size of the training dataset for training will be prohibitively large.

Thus, our recognition model in this paper adopts few-shot learning techniques [2, 4, 12, 23, 28], making it the first on few-shot human atomic action recognition. Our few-shot model, combined with a global alignment model, performs matching using explicit 4D skeletal representation. Exploiting geometry inherent in 4D human skeletons avoids training complex models to learn proper encoding in the embedded space, which may still not produce better alignment than using explicit alignment (Figure 2) and thus hampering recognition. Our experiments validate that the proposed few-shot model working in tandem with a global alignment model produces better recognition results with fewer training examples for each action. In summary, our contributions consist of:

1. HAA4D, a human atomic action dataset where all 4D skeletons are globally aligned. HAA4D complements existing prominent 3D+T human action datasets such as NTU-RGB+D [25, 31], and Kinetics-skeleton [19]; HAA4D contains 3390 samples of human actions in the wild with 300 different kinds of activities. The samples for each human action range from 2 to 20, each provided with RGB frames and their corresponding 3D skeletons;

2. introducing an alignment network for predicting orthographic camera poses in the train/test samples, where all 4D skeletons are aligned in the same camera space each facing the negative $z$-direction. This allows for better recognition results with small number of training samples compared to ST-GCN[40], Shift-GCN[8], and SGN[43];

3. the first few-shot baseline for 4D human (atomic) action recognition that produces results comparable to or better than state-of-the-art techniques on unseen classes using a small number of training samples.

2. Related Works

2.1. Human Video Action Datasets

In skeletal human action recognition, several datasets are often used as benchmarks, see Table 1. NTU-RGB+D [31] and NTU-RGB+D 120 [25] were collected under laboratory settings that cover actions in three main categories: daily actions, mutual actions, and medical conditions. These datasets comprise of RGB videos, depth map sequences, 3D skeletal data, and infrared videos for each sample. There are other RGB+D datasets that were also captured in laboratory settings, such as UWA3D Multiview II [30] which contains 30 various activities, and SYSU 3D HOI [16] which provides 12 activities with 40 different participants focusing more on human-object interaction. Recently, Ego4D [14] has been released which contains massive-scale egocentric videos and provides long sequences of actions in real-life scenarios, where parts are given with the corresponding 3D meshes. However, although having a massive number of samples per action, these datasets are often not sufficiently diversified, subject to the drawback of not covering a wide range of different actions.

Kinetics-skeleton, on the other hand, contains more diversified human actions. This skeleton dataset was built on top of Kinetics [19], a video action datasets collected from YouTube with more than 300,000 video clips covering 400
Datasets Samples Actions Views Modalities Year
UW A3D Multiview II [30] 1075 30 5 RGB+D+3DJoints 2015
NTU RGB+D [31] 56,880 60 80 RGB+D+IR+3DJoints 2016
SYSU 3DHOI [16] 480 12 1 RGB+D+3DJoints 2017
Kinetics 400 [19] 306,245 400 - RGB 2017
NTU RGB+D 120 [25] 114,480 120 155 RGB+D+IR+3DJoints 2019
Kinetics-700-2020 [32] 633,728 700 - RGB 2020
Ego4D [14] 3,025 110 - RGB+3DMeshes+Audio 2021
HAA4D 3,390 300 2/20 RGB+3DJoints 2021

Table 1. Comparison between HAA4D and other public datasets for human action recognition. With fewer samples than many existing datasets, HAA4D provides class-balanced and diversified actions with 3D+T or 4D skeleton data useful for few-shot action learning.

action classes. OpenPose [7] was used to extract 2D human skeletons. Together with EvoSkeleton [24] used to lift the 2D key-points to 3D skeletons, we can extract the 3D skeletal data from the images. While Kinetics-skeleton introduces more variety of human poses to the human skeletal action recognition domain, the correctness of the 3D skeletons generated may be questionable, as the 2D joints prediction network can fail if parts of the human body are outside of the image frame.

To ameliorate the problem of existing datasets problem, namely, lack of pose diversity and low ground-truth accuracy, we introduce HAA4D which contains 300 human action classes, each with 20 examples with accurately annotated 2D joints position.

2.2. Video-Based and Few-Shot Action Recognition

For more recent examples on video-based action recognition, C3D [33] performs 3-dimensional convolution on the input image sequence to extract spatio-temporal features. In [34] an architecture is proposed for first processing each image using 2D convolution, and then using a bidirectional LSTM network to learn temporal information. TSN [37] divides a video into several segments and selects snippets to pass to a spatial stream ConvNets and a temporal stream ConvNets. Video-level prediction is then derived from the consensus of the snippets. However, all of these models contain an excessive number of learnable parameters that require training on large-scale datasets, which can fail when training samples are few and expensive to obtain. Thus, more works have focused on few-shot video action classification.

In [29] the authors proposed to learn to generate class embeddings using a Word2Vec [27] model. NGM [15] introduces a graph matching metric on a graphical representation of a video. Dense dilated network [39] uses a dilated CNN network on videos. In contrast to these works that neglect the temporal dimension of videos, TARN [4] introduces temporal attention for temporal alignment of videos. A similar approach is used in [41] by adopting temporal attention as a temporal form of self-supervision. Our work is closest in spirit to OTAM [5] which uses an ordered temporal alignment module, inspired from dynamic time warping (DTW) [1]. However, instead of focusing only on temporal alignment, HAA4D explicitly aligns along 3D+T or 4D dimensions since similar actions, after calibration removing the view and scale variation, share similar joint distribution and moving sequences.

However, all of the aforementioned approaches directly train a video classification model using RGB-frames, which poses a deep neural network a great challenge in relating video frames as they can vary widely in background and illumination. In this paper, we use skeleton data extracted from the videos for better spatio-temporal alignment.

2.3. Skeleton-Based Action Recognition

Given the aforementioned advantages, skeleton-based action recognition has been a popular topic in computer vision that aims to classify actions using skeleton joint information. Conventional methods rely on hand-crafted features to capture dynamic movement of human joints [17, 35, 36]. With emergence of deep-learning, earlier related works represent joint coordinates as a vector, and use recurrent neural network for action classification [10, 26, 42], or represent skeleton sequence as a pseudo-image as input to a CNN-based network [20, 21, 22]. Recently, ST-GCN in [40], which consists of spatial graph convolution layers and temporal convolution layers, uses graph representation of skeleton sequence. ST-GCN has been extended in [8] by incorporating shift convolution [18, 38, 44] on a graph CNN, which achieves better performance with a lighter network. In this paper, we use Shift-GCN [8] as a backbone for our baseline model to compare the efficacy of our globally aligned 4D skeletons.

3. HAA4D

This section will introduce details and structure of our HAA4D dataset, its expandability, and the evaluation criteria. Figure 3 and 4 shows some examples of our HAA4D dataset. Full details of HAA4D are available in the supplementary materials.
HAA4D consists of 3,300 real-life RGB videos of specific human action classes with the corresponding 2D joints and 3D human skeletons. The skeleton topology follows the one in Evoskeleton [24] which consists of 17 joints which are quite sufficient in representing a wide range of human body and part movements. The dataset has 300 action classes that are divided into two categories: primary classes and additional classes. Primary classes have 20 samples per class, while additional classes contain two examples per class. Recall the main advantage of 4D skeleton data is its viewpoint invariance with a few as one 3D skeleton per frame, given they are globally aligned with camera poses available which are estimated using our camera prediction model.

For primary classes, videos 0–9 are used for training, and the remaining are for evaluation. Actions in the additional classes are dedicated to one-shot learning that evaluates the performance of the proposed model to differentiate the action when very limited data are presented. For example, this paper trains the alignment model on the first ten examples of human movements in the primary classes with data augmentation techniques to be detailed. The alignment model is then tested with videos in the evaluation sets of the primary classes for cross-view evaluation, and with examples in the additional classes for cross-action assessment.

### 3.2. Mean Average Precision (mAP)

We calculated the mean average precision (mAP) of the whole dataset and each primary class using the bounding boxes of the predicted skeletons generated from AlphaPose [11] and our human-labeled ground truth skeletons to calculate the IoU. The threshold of the IoU is set to 0.5. The overall mAP of the HAA4D dataset is 63.71. However, we observed a significant variance between the mAP of each class. For example, the mAP of the action ‘abseiling’ is 92.08 while the mAP of ‘applying_cream’ is 1.28. One reason that caused the difference is that most images captured contain only the upper half of the body for activities like ‘applying_cream’. This shows the insufficiency of state-of-the-art pose prediction networks like AlphaPose in predicting joints that are out of boundary; therefore, making our perfectly labeled ground truth more valuable. Although bounding boxes cannot effectively represent each joint’s position accuracy, we will provide part-level bounding annotation for more accurate evaluation. The mAP details for each action class will be included in the supplemental material.

### 3.3. Annotation Tool

To help expand the dataset to more actions and classes, we have developed a simple interactive annotation tool that supports faster user labeling (Figure 5). Similar to Kinetics-skeleton use of OpenPose [7] to predict positions of 2D joints, the user can load preliminary predictions from AlphaPose [11] and can correct intermediate frames where the network prediction is not precise. Observing that human movement when viewed in a short period is relatively linear, the annotation tool supports linear interpolation so that the user can avoid frame-by-frame processing whenever possible. The predicted 3D skeleton is also shown alongside in
the interface for the user to monitor the generated ground truth in real time. The annotation tool covers the end-to-end generation of 3D human skeletons from images, which is user-friendly and efficient in annotation, thus making this HAA4D dataset easily expandable. We hope that by providing this annotation tool, the dataset can become more comprehensive in the future to cover more human action poses, contributing to research in few-shot human pose estimation and action recognition that utilizes 3D+T data.

3.4. Globally Aligned Skeletons

Since examples in HAA500 were shot in the wild, each action sample has different viewpoints and scales, making the predicted 3D skeletons in various coordinate systems. Therefore, in HAA4D, we apply the following to standardize the 3D generated skeletons.

First, all skeletons are centered such that their lower-spine joint is placed at the origin. Second, because EvoSkeleton [24] generates 3D skeletons frame-by-frame, it does not guarantee equal bone lengths throughout the video. Hence, to ensure the human skeleton’s uniformity in the action video, we use bone lengths in the first frame as the reference to replace the lengths of the rest frames. We also scale all the bones such that their l2-norm equals to one, i.e., $b_i^f$ represents bone $i$ in the $f$-th skeleton frame and $B$ is the set of bones in the skeleton topology.

Then, we provide globally aligned skeletons (to parts of the dataset) by manually rotating them so that the human faces the negative $z$-direction at the start of each action. We choose the action to face in the negative $z$-direction because most people when taking photos will look into the camera. We calculate the alignment using only the first frame and apply the rotation to the rest of the sequence as a frame-by-frame alignment will break the relative movement of the original action. The calibration makes our data view-invariant by bringing all skeletons to the same space. We will argue in the next section that eliminating the rotation and scale factors help us better compare the skeleton movement in different actions by directly using the adjusted 3D joints position. The manually corrected, globally aligned skeletons are also used to train the alignment model, which takes any actions, regardless of their view angles and action type, and transforms them to the global coordinate system.
4. Few-Shot Action Recognition Model

In this section, we describe our few-shot skeleton-based action recognition model. Figure 6 shows the overall architecture which consists of two modules: the global alignment module (with data augmentation) and the sequence matching module.

Following [5, 37], given a video example, we divide it into $t_n$ segments and randomly select a frame within each segment. We randomly selected one frame from each segment to introduce more randomness to each input sequence. As we work on atomic actions, given the same action type, the start and end will match. However, each person may act at a different pace, and thus, we do not hardcode any frame to represent each segment.

We then concatenate the $t_n$ skeletons to represent the video, denoted by $S$. This allows each video to be represented by the same number of frames. Similarly, the support sets are constructed by randomly selecting $s_n$ samples from each $w_n$ class, including the query’s. The global alignment model transforms both the query and support into the globally aligned space before action recognition, by comparing the sequence matching results with the support sets.

4.1. Global Alignment Model (GAM)

As query and support videos can be captured in-the-wild, we propose the global alignment model to bring all inputs to the same space, such that actions face in the negative $z$-direction at the beginning. The camera angle will also be available after our global alignment.

To prepare or synthesize training data, since we have the ground truth globally aligned skeletons as described in section 3.4, we perform data augmentation by uniformly sampling different camera views from vertices on an $n$-frequency icosahedron (i.e., subdivided icosahedron incident on a unit sphere as shown in Figure 1). Our globally aligned skeletons are viewed from the default orthographic camera $C_D$ located at $(0, 0, -1)$. By simplifying the camera to have no self-rotation, we only need to predict two rotation parameters, namely, azimuth and altitude angles, denoted by $\beta$ ($\theta$ and $\phi$) with respect to the default camera position $(0, 0, -1)$. We obtain $\theta$ and $\phi$ by:

$$\theta = \pi - \arctan\left(\frac{C_x}{C_z}\right), \quad \theta = \begin{cases} \theta - 2\pi, & \text{if } \theta > \pi \\ \theta, & \text{otherwise} \end{cases}$$  \quad (1)

$$\phi = \arctan\left(\frac{C_y}{\sqrt{(C_x)^2 + (C_z)^2}}\right)$$  \quad (2)

We then use Rodrigues rotation to obtain the rotation matrix $R = R_2 \cdot R_1$, where:

$$\text{Rodrigues}(\hat{n}, \alpha) = I_3 + \sin \alpha K + (1 - \cos \alpha)K^2,$$

where $K = \text{Skew}(\hat{n})$  \quad (3)

We first rotate in $\theta$ using:

$$R_1 = \text{Rodrigues}(\hat{n}_1, \theta), \quad \hat{n}_1 = \langle 0, -1, 0 \rangle$$  \quad (4)

Then rotate in $\phi$: let $\vec{v}_1 = \langle 0, 1, 0 \rangle$, $\vec{v}_2 = R_1 \cdot C_D$, then

$$R_2 = \text{Rodrigues}(\hat{n}_2, |\phi|), \text{ where } \hat{n}_2 = \begin{cases} \vec{v}_2 \times \vec{v}_1, & \text{if } \phi > 0 \\ \vec{v}_1 \times \vec{v}_2, & \text{otherwise} \end{cases}$$  \quad (5)
We obtain the augmented view skeleton $S_A$ by rotating the globally aligned skeletons $S_G$:

$$S_A = R^{-1} \cdot S_G,$$

where $R = R_2 \cdot R_1$.

We sample data from 92 camera angles on a 3-frequency subdivision icosahedron, trained on 73 views, and test on the rest 19 views. We employ rotation loss and reconstruction loss defined as follows:

$$L_{rot} = \text{MSE}(\cos \beta_{pr}, \cos \beta_{gt}) + \text{MSE}(\sin \beta_{pr}, \sin \beta_{gt})$$

(7)

$$L_{rec} = \text{RMSE}(S_{pr}, S_A),$$

where $S_{pr} = (R(\beta_{pr}))^{-1} \cdot S_G$.

The final loss for our global alignment model is

$$L = w_1 L_{rot} + w_2 L_{rec},$$

(9)

where we set $w_1$ to 10 and $w_2$ to 1.

4.2. Sequence Matching

After transforming all input skeletons to the same global space, where we denote each transformed skeleton as $S$, we encode the 3D positions of the joints $f_{pos}(S)$ and their respective trajectory $f_{tra}(S)$ as embeddings for sequence matching. The encoding functions are defined as follows:

$$f_{pos}(S) = S^1 \oplus S^2 \oplus \cdots \oplus S^{t_n}$$

(10)

$$f_{tra}(S) = 1 \oplus (S^2 - S^1) \oplus \cdots \oplus (S^{t_n} - S^{t_n-1})$$

(11)

$$f_{skel}(S) = f_{pos}(S) \oplus f_{tra}(S)$$

(12)

where $\oplus$ denotes concatenation between the two frames. Inspired by OTAM [5] and dynamic time warping [1], we use the matching score between the explicit query $f_{skel}(S_q)$ and supports $f_{skel}(S_s)$ in embedding to determine the action class of the given query. We generate the distance matrix $D = 1 - \cosine_similarity(f_{skel}(S_q), f_{skel}(S_s))$. The score of the query and the support is represented by $- \sum_d d_{path} d$, where $path$ is a relaxed-minimum path of $D$ defined in [5]. The final score of a query and a support class is the average of scores for each $k$ support data of the class. The top final score corresponds to the action class of the query video.

5. Experiments

5.1. Skeletal Representation

Implicit vs Explicit Without representative skeletal-based few-shot action recognition models, we create a baseline model to evaluate the performance. The baseline model uses Shift-GCN [8] as the backbone to encode input skeletons in an embedded space and perform the temporal alignment. The Shift-GCN model is first pre-trained with the embedding cosine similarity loss and the skeleton reconstruction loss. To simulate the few sample situations in the dataset, we randomly select ten samples from each NTU-RGB+D [31] and for training, and use the other ten examples for testing. The encoded skeleton in the embedded space is the implicit representation of the skeleton sequence used for sequence matching.

2D vs 3D Here we compare the effectiveness between the 2D and 3D skeletons. The most notable difference between using 2D and 3D skeletons is that for 2D skeletons, we cannot perform data augmentation as rotation cannot be done in the 2D space without breaking the original skeleton structure. We experiment on both the baseline model and other proposed architectures. In the baseline model, we follow the same approach to encode the 2D skeletons to an implicit representation in the embedded space. However, for testing on our model, we directly feed 2D skeletons for sequence matching without the proposed 3D global alignment.

Results Table 3 shows the effectiveness of using different representations of human skeleton sequences for action recognition. The baseline model utilizes implicit representation, while our model uses explicit information. We test both models with 2D and 3D as input skeletons and demonstrate that explicit 3D representation, i.e., globally aligned information, produces the best accuracies in action recognition.

5.2. Ablation Studies

In this section, we conduct ablation studies to examine the effect of individual part of our model. We tested on three different factors: the number of sampled frames, the sequence matching approach, and the explicit information used as encodings. We monitor the results on both HAA4D and NTU-RGB+D.

**Sampling Rate** For each human skeleton action sequence, we divide them into several segments and select one skeleton from each to represent the segment. This sampling technique allows all the action sequences to be in uniform length while increasing the difference between the neighboring frames. We test on different numbers of frames to represent the sequence, as oversampling can lead to redundancy and unnecessary computational load, while undersampling can lead to temporal aliasing. We padded the
last frame to extend the input sequence for input with fewer frames than the segment size.

Sequence Matching Our action recognition model determines the action by comparing the sequence matching score between the query set and support sets. We experiment the following matching techniques:

- **Mean**, **Regular DTW**, and **OTAM** [1, 5]. The Mean approach for sequence matching directly computes the average between the two distance matrix $D$ and thus neglects the temporal ordering. The other two methods consider temporal ordering. Regular DTW assumes the two sequence matches at the start and end, which allows movement in the direction of $\rightarrow$, and $\downarrow$ when computing the aligned path in the distance matrix. OTAM on the other hand assumes the alignment can start and end from any time in the sequence, thus only allowing $\rightarrow$ and $\downarrow$ to ensure that all the alignment paths have equal length.

Explicit Encodings In addition to directly using the aligned 3D coordinates ($0^{th}$ order) of the skeleton sequence as our explicit embeddings, we evaluate the effect of using different combinations between the $0^{th}$, $1^{st}$, and $2^{nd}$ order representations. The $1^{st}$ order of the skeleton sequence consists of the joints movement throughout temporal frames. The $2^{nd}$ order representation includes the curvature of the joint path.

Results From our experimental results tabulated in Table 4, the 32-frame sampling outperforms others in both HAA4D and NTU-RGB-D datasets. In addition, we observe that techniques that consider temporal ordering, such as regular DTW and OTAM, tend to produce better accuracy in action prediction for the sequence matching methods. Notably, the actions in HAA4D are atomic, which are diverse including atomic motions that form more complex actions, where the actions in the same class match at the beginning as well as the end of the sequence. This feature benefits the regular DTW to obtain more reliable results.

|                | NTU-RGB+D xview [31] | HAA4D |
|----------------|----------------------|-------|
|                | Sway 1shot | Sway 5shot | Sway 1shot | Sway 5shot |
| Mean           |           |           |           |           |
| - $t_n = 8$    | 47.5      | 51.0      | 45.9      | 54.8      |
| - $t_n = 32$   | 51.8      | 59.8      | 47.9      | 56.5      |
| - $t_n = \text{all}$ | 52.0 | 60.5  | 47.5 | 53.2 |

|                |                  |       |       |
|----------------|------------------|-------|
|                | Regular DTW      |       |       |
| - $t_n = 8$    | 58.0             | 69.8  | 52.4  | 62.3 |
| - $t_n = 32$   | 59.3             | 71.3  | 53.5  | 62.7 |
| - $t_n = \text{all}$ | 59.1 | 69.0 | 53.4 | 62.5 |

|                |                  |       |       |
|----------------|------------------|-------|
|                | OTAM             |       |       |
| - $t_n = 8$    | 56.5             | 67.8  | 51.9  | 62.0 |
| - $t_n = 32$   | 57.8             | 70.0  | 52.1  | 62.3 |
| - $t_n = \text{all}$ | -    | -    | -    | -    |

Table 4. Performance of few-shot action classification under different sequence matching methods and sampling rates. We do not consider $t_n = \text{all}$ for OTAM as it constrains the distant matrix to be squared and having equal length paths.

Table 5 shows that by using both the globally aligned position and the joints temporal movement, the model produces the best performance.

5.3. Comparison with State-of-the-Arts

This section evaluates state-of-the-art skeleton-based action recognition models on our HAA4D dataset, and compares the results of the testing on other public datasets. Although few-shot learning has been used on action recognition [5], few-shot learning has been conducted via the 3D or 3D+T skeleton-based method. In addition, one of our claims is the use of few-shot learning on 3D skeleton human action recognition to improve the prediction accuracy in sparse training training data. We compare our result with the state-of-the-art supervised approaches to support our claim.

The skeletons of HAA4D are processed according to the description in section 3.4, which centers the origin and constrains all skeletons to have equal bone length throughout the video. The inputs of ST-GCN [40] and Shift-GCN [8] follow the shape (batch-size $\times$ channels $\times$ frames $\times$ joints $\times$ people). We let the channels be the $(X, Y, Z)$ of coordinates of the 3D skeletons and pad zeros to the second person in people dimensions, as HAA4D targets only single-person action. SGN [43] splits a multi-person frame into multiple frames, in which each frame contains only a single person. This model also requires segmenting the skeleton sequence into 20 clips, where one frame is randomly sampled from each segment to form the new sequence. We follow the same data processing techniques to evaluate our HAA4D.

Table 6 shows that all models suffer from the lack of accuracy in Kinetics-skeleton. One possible reason is that the skeletons are directly extracted from the Kinetics using OpenPose [7] without correction done on the corrupted skeletons, where the 2D predictions are not sufficiently accurate to be lifted to the corresponding 3D skeleton. On the other hand, our HAA4D is a curated dataset with precise 2D joints location; even out-of-frame joints are manually completed reasonably. Thus the ground truth 3D skeleton is more reliable. This leads to better accuracy for the state-of-the-art models on our dataset. However, noting that HAA4D is designed for few-shot action recognition and due to the limited size of HAA4D, there is still a gap in prediction precision between HAA4D and NTU-RGB+D [31].
Table 6. Performance of state-of-the-art skeleton-based action recognition models on different datasets.

|                     | ST-GCN [40] | Shift-GCN [8] | SGN [43] |
|---------------------|-------------|---------------|----------|
| NTU-RGB+D [31]     |             |               |          |
| - xview             | 88.8        | 95.1          | 94.5     |
| - xsub              | 81.6        | 87.8          | 89.0     |
| Kinetics-skeleton [19] | 31.6        | 17.7          | 20.8     |
| Ours (HAA4D)        | 21.2        | 53.0          | 53.3     |

Figure 7. Accuracy respect to each action classes

6. Discussion

6.1. Substantial need for the clean HAA4D dataset

One may concerned that “clean” dataset is not practical in real-life scenarios. However, current 2D pose prediction networks are still far from the accuracy of hand-corrected labeling. With much effort, introducing this cleaner dataset can advance this area of studies as all images collected in HAA4D are “in the wild” with more accurate ground truth 2D skeletons. Moreover, we observed several actions in NTU RGB+D 120 [25] and Kinetics-skeleton [19] are corrupted, including wrong subjects or unreasonable joint movements, that humans cannot even recognize. Our highly diversified HAA4D does not have these problems to distract learning. Also, having a clean dataset, we can add and control noises to generate false actions and even train networks to identify these errors.

6.2. Evaluation on failure cases

Figure 7 shows the worst 5 classes for our action recognition model are yawning (0%), water_skiing (2%), alligator_wrestling (3%), balancebeam_jump (3%), yoga_gate (5%). Actions such as yawning mainly involve mouth movements, which is difficult to reflect using skeletons. The model is subject to cases that lack significant movements in parts of the skeleton. Similar cases can be found in water_skiing and alligator_wrestling. The model also obtains low accuracy when the actions contain significant variances and lack consistency. For actions like balancebeam_jump, different dancers have unique styles and thus poses. It is the balance beam (interacting object) that helps humans identify the action even though each example looks completely different.

7. Conclusion

This paper proposes a new 4D dataset HAA4D which consists of more than 3300 RGB videos in 300 human atomic action classes. HAA4D is clean, diverse, class-balanced where each class is viewpoint-balanced with the use of 3D+T or 4D skeletons. All training and testing 3D skeletons in HAA4D are globally aligned. Such alignment makes matching skeletons more stable, and thus with fewer training samples per class needed for action recognition leveraging full 3D+T information, in stark contrast to the 2D counterparts where massive amount is required to cover adequate viewpoints.

Given the high diversity and skeletal alignment in HAA4D, we construct the first baseline few-shot 4D human atomic action recognition network, which produces comparable or higher performance than relevant state-of-the-art techniques, using the same small number of training samples of unseen classes. Through ablation studies, we have verified the advantages of 3D skeletons over 2D counterparts. We have also found that existing 3D skeletal action datasets (e.g., NTU-RGB+D [31] and Kinetics-skeleton [19]) significantly benefit from the proposed explicit alignment in the few-shot setting. With its high scalability where only a small number of 3D+T skeletons to be annotated for class extension using our annotation tool, and 3D+T global alignment contributed by this paper, we hope HAA4D will spawn fruitful future works on skeletal human action recognition. Dataset, codes, models and tools will be released upon publication.

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HAA4D: Few-Shot Human Atomic Action Recognition via 3D Spatio-Temporal Skeletal Alignment (Supplementary Material)

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1. Action Classes
This section provides the complete list of actions contained in our HAA4D dataset. The dataset can be characterized in two parts: primary classes and additional classes, where the former contains 155 actions and the latter contains 145 actions.

1.1. Primary Classes
Table 1 shows actions in the primary classes. Primary classes are actions that contain 20 samples per class. We also include the mean average precision of each class.

1.2. Additional Classes
Table 2 shows actions in the primary classes. Additional classes are actions that contain two samples per class, which are dedicated to one-shot learning. It helps evaluate the performance of the proposed model to differentiate the action when very limited data are presented.

2. Globally Aligned Skeletons
Globally aligned skeletons are skeletons that we manually rotated so that all the samples are facing the negative z-direction at the start of the action. This adjustment eliminates view variances so that all actions to be performed in the same coordinates space. There are 40 different actions in the primary classes containing globally aligned skeletons. We will provide the list in Table 3.

3. HAA4D Evaluation Benchmark
For actions with 20 examples per class, video indexes 0 to 9 are used for training, while videos from 10 to 19 are used for testing. We perform data augmentation on the first ten samples, and among all, videos 8 and 9 are used for validation. For actions that contain only two samples, the one with a smaller index serves as the query, and the other serves as the support.

4. Training Configuration
For training the global alignment network, we perform data augmentation by sampling camera views from a 3-frequency subdivision icosahedron. This gives us 92 additional training samples per example. Since there are 400 examples in HAA4D that are provided with globally aligned skeletons, with the help of data augmentation, we have 36,800 examples of training our global alignment network. We split all the data into training and validated with a ratio of 0.8 under two settings: cross-views and cross-actions. We select 73 views on the icosahedron sphere for cross-views and test the rest 19 views to ensure that our network generates predictions decently while encountering unseen views. We also trained our network on cross-views, i.e., 32 out of the 40 classes, to secure that the model is used to generalize different actions. Here are our training environment and configurations in more details:

- GPU: GeForce GTX TITAN X and GeForce GTX 1080 Ti
- Epochs: 300
- Batch size: 64
- Optimizer: Adam (starting learning rate: 1e-4, weight decay rate: 1e-6)
- Loss weight: 10:1 (rotation loss : reconstruction loss)

5. Limitations
Our dataset contains only a few samples per class, which is intended for enhancing its scalability and extension (e.g. 1 for train and 1 for test, see our 5-way 1-shot experiments in main paper), as unlike the 2D counterparts, HAA4D’s 3D+T skeletons have more degree of freedom making meaningful data augmentation easy for training on large datasets. In our case, we sample the 3D skeletons from different viewpoints and rotate the skeletons accordingly. We can also use mirroring or combining skeleton samples in the same class to introduce more variation to the dataset. Unlike 2D skeletons that can only have one rotation parameter, the properties of 3D skeletons help better perform data augmentation without breaking the original skeleton structure.

Our few-shot skeleton-based action recognition architecture currently supports only single-person actions. To
accommodate actions involving more than one person, we can use a similar technique as ST-GCN, which utilizes the two skeletons with the highest confidence score in the sequences. For actions that have only one subject, we assume them to be all zeros. With this, we can modify our architecture so that instead of having the shape of \((n, \text{ways}, n, \text{encodings})\) for explicit skeletons encodings, we add one additional dimension so that the skeletons are in the shape of \((n, \text{ways}, n, \text{s}, n, \text{s})\).

We then calculate the distance, respectively. Since there are \(n\) ways, \(n\) segments, \(n\) encodings for explicit skeletons encodings, we add one additional dimension so that the skeletons are in the shape of \((n, \text{ways}, n, \text{s}, n, \text{s})\).

We then calculate the distance, respectively. Since there are \(n\) ways, \(n\) segments, \(n\) encodings for explicit skeletons encodings, we add one additional dimension so that the skeletons are in the shape of \((n, \text{ways}, n, \text{s}, n, \text{s})\).
the minimum distance between the possible combination
d(q,s1, s,s1) + d(q,s2, s,s2), d(q,s1, s,s2) + d(q,s2, s,s1).
Therefore, we can make this adjustment for multi-person
interaction, and the rest of the architecture can remain
unchanged.

Table 3. Actions the contain globally aligned skeletons.