Int&Int: A Two-Pathway Network for Skeleton-Based Action Recognition

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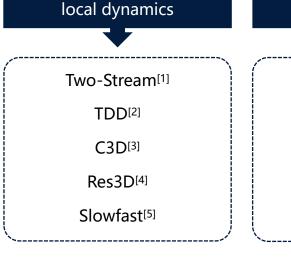
Research Status

Our Framework

Contributions

Research Status

- Mainly focus on the local or global information alone.
- Cannot focus on both the intensity and the integrity.



global information

TSN^[6]

TRN^[7]

TLE^[8]

ActionVLAD[9]

LTC[10]

Timeception^[11]

Methods

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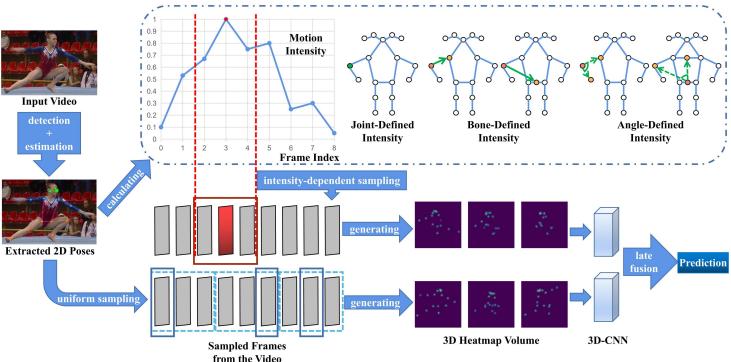
Research Status

Our Framework

Contributions

Our Framework

We propose a two-pathway *Int&Int* network(*Intensity&Integrity*) for skeleton-based action recognition to satisfy both aspects.



Research Status

Our Framework

Contributions

Contributions

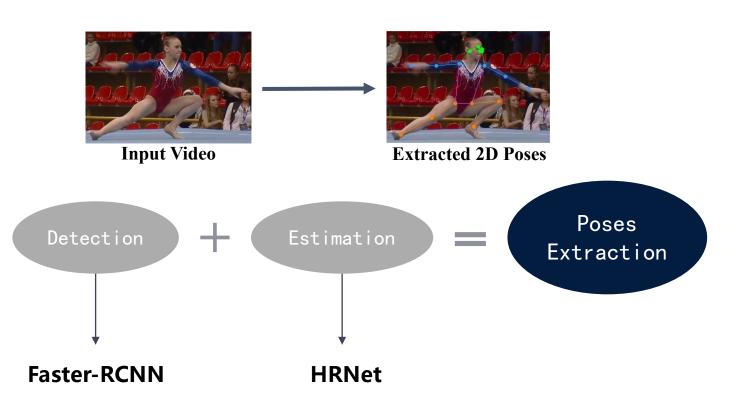
- 1) A two-pathway *Int&Int* framework is proposed which concentrates on the local and global information of the action, where there is great complementarity in the two pathways.
- 2) A clip whose position depends on the most intense frame is sampled as for *Intensity* pathway, where there is a clear physical meaning in the sampling process.
- 3) The motion intensity of each frame is defined in 3 + 2 ways containing different semantic information.

Poses Extraction

Two Pathways

Late Fusion

Poses Extraction

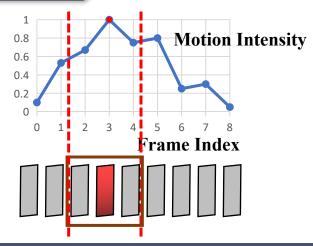


Poses Extraction

Two Pathways

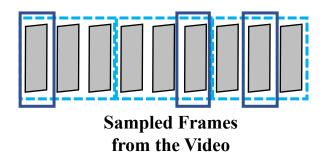
Late Fusion

Two Pathways



Intensity-dependent sampling in Intensity pathway

- Motion features are **non-uniformly** distributed along the time axis.
- Motions with large amplitude and quick changes have more actionspecific semantic information for identifying the action intuitively.
- Intensity-dependent sampling can avoid missing the salient features:
- 1) extracted 2D poses --> motion intensity of each frame in the video
- 2) select the largest element corresponding to the most intense frame
- 3) sample a clip composed of consecutive frames around the most intense frame



Uniform sampling in Integrity pathway

- Maintain the global information of video:
- 1) one input video is divided into n segments of equal length (n frames to sample)
- 2) one frame is randomly selected from each segment

Poses Extraction

Two Pathways

Late Fusion

Two Pathways

3D heatmap volumes

• We generate a joint heatmap J by composing K gaussian maps centered at every joint, which are based on joint coordinates and confidence: $(i-x, i)^2 + (i-y, i)^2$

 $J_{kij} = e^{-\frac{(i-x_k)^2 + (j-y_k)^2}{2*\sigma^2}} * c_k$







3D Heatmap Volume

- In the multi-person case of one frame, the k-th gaussian maps of all persons are directly accumulated.
- For all sampled T frames, we stack all heatmaps along the temporal dimension to obtain a 3D heatmap volume of size $K \times T \times H \times W$, where K is the number of joints, H and W are the height and width of the frame.
- Subjects-centered cropping is also adopted to reduce the redundancy of 3D heatmap volumes.

3D-CNN of two pathways

- We use the pseudo heatmap volumes as the input.
- $T \times S^2$, C denote the dimensions of kernels for temporal, spatial, channel sizes.
- We choose SlowOnly, obtained by inflating the ResNet layers in the last two stages from 2D to 3D, to instantiate the backbone.
- Both pathways have the same architecture, and they train the individual losses respectively.

Stage	Pathway	Output Size $T \times S^2$		
Data Layer	$32,4^{2}$	32×56^{2}		
Stem Layer	Conv $1 \times 7^2, 32$ Stride $1, 1^2$	32×56^2		
ResNet3	$\begin{bmatrix} 1 \times 1^2, 32 \\ 1 \times 3^2, 32 \\ 1 \times 1^2, 128 \end{bmatrix} \times 4$	32×28^2		
ResNet4	$\begin{bmatrix} 3 \times 1^2, 64 \\ 1 \times 3^2, 64 \\ 1 \times 1^2, 256 \end{bmatrix} \times 6$	32×14^2		
ResNet5	$ \begin{array}{ c c c c c c } \hline 3 \times 1^2, 128 \\ 1 \times 3^2, 128 \\ 1 \times 1^2, 512 \end{array} \times 3 $	32×7^2		
GAP	GlobalAveragePooling	#Classes		

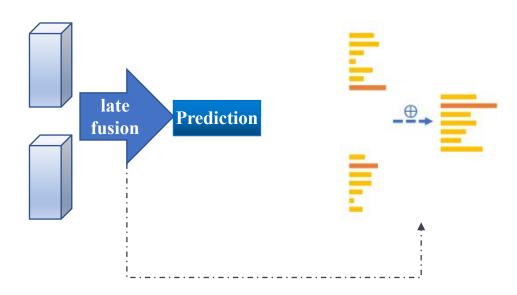
Poses Extraction

Two Pathways

Late Fusion

Late Fusion

The softmax scores of the two pathways are added to obtain the fused score and predict the corresponding action category.



Joint-Defined

Bone-Defined

Angle-Defined

JBA-Defined

Joint-Defined

The final value of joint-defined motion intensity is the arithmetic average of values calculated and normalized in two ways respectively.

1) Using corresponding position in the next frame

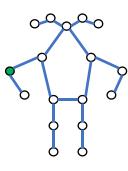
- calculate the distance between corresponding positions of adjacent frames
- J_{ij}^t means the position of the *i*-th keypoint of the *j*-th person in the *t*-th frame. K is the number of keypoints of the skeleton. M is the number of people in the frame:

$$SV_{Jnf}^{t} = \frac{\sum_{j=0}^{M-1} \sum_{i=0}^{K-1} |J_{ij}^{t+1} - J_{ij}^{t}|}{M}$$

2) Using average position across the frames

- calculate the distance between the *i*-th keypoint in the *t*-th frame and the average position of the keypoint
- \bar{J}_{ij} means the average position of the *i*-th keypoint of the *j*-th person across all frames:

$$SV_{Jap}^{t} = \frac{\sum_{j=0}^{M-1} \sum_{i=0}^{K-1} |J_{ij}^{t} - \bar{J}_{ij}|}{M}$$



Joint-Defined

Bone-Defined

Angle-Defined

JBA-Defined

Bone-Defined

The final value of bone-defined motion intensity is the arithmetic average of 2×2 values (2 ways to calculate $\times 2$ types of bone features) which have been normalized.

1) Using corresponding bone-vector in the next frame

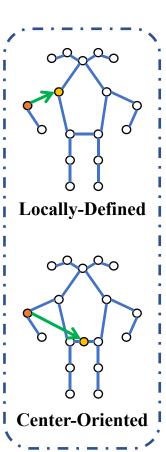
- calculate the angle between bone-vector $B^t = \overrightarrow{J^t v^t}$ and $B^{t+1} = \overrightarrow{J^{t+1} v^{t+1}}$
- B_{ij}^t means the *i*-th bone-vector of the *j*-th person in the *t*-th frame:

$$SV_{Bnf}^{t} = \frac{\sum_{j=0}^{M-1} \sum_{i=0}^{K-1} \left(1 - \frac{B_{ij}^{t} \cdot B_{ij}^{t+1}}{|B_{ij}^{t}| |B_{ij}^{t+1}|}\right)}{M}$$

2) Using average bone-vector across the frames

- calculate the angle between bone-vector $B^t=\overrightarrow{J^tv^t}$ and $\bar{B}=\overrightarrow{\bar{J}v}$
- \bar{B}_{ij} means the *i*-th average bone-vector of the *j*-th person across all frames:

$$SV_{Bap}^{t} = \frac{\sum_{j=0}^{M-1} \sum_{i=0}^{K-1} \left(1 - \frac{B_{ij}^{t} \cdot \bar{B}_{ij}}{|B_{ij}^{t}||\bar{B}_{ij}|}\right)}{M}$$



Joint-Defined

Bone-Defined

Angle-Defined

JBA-Defined

Angle-Defined

The final value of angle-defined motion intensity is the arithmetic average of 2×2 values (2 ways to calculate×2 types of angular features) which have been normalized.

1) Using corresponding angle in the next frame

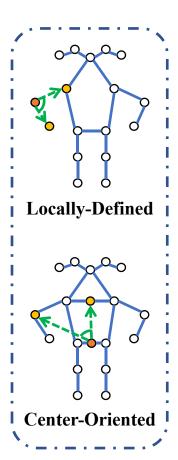
- calculate the difference between corresponding angles of adjacent frames
- $A_{ij}^t=1-\cos\theta_{ij}^t$ where θ_{ij}^t means the *i*-th angle of the *j*-th person in the *t*-th frame:

$$SV_{Anf}^{t} = \frac{\sum_{j=0}^{M-1} \sum_{i=0}^{K-1} |A_{ij}^{t+1} - A_{ij}^{t}|}{M}$$

2) Using average angle across the frames

- calculate the difference between the *i*-th angle in the *t*-th frame and average angle
- $\bar{A}_{ij}=1-\cos\bar{\theta}_{ij}$ where $\bar{\theta}_{ij}$ means the *i*-th average angle of the *j*-th person across all frames:

$$SV_{Aap}^{t} = \frac{\sum_{j=0}^{M-1} \sum_{i=0}^{K-1} |A_{ij}^{t} - \bar{A}_{ij}|}{M}$$



Joint-Defined

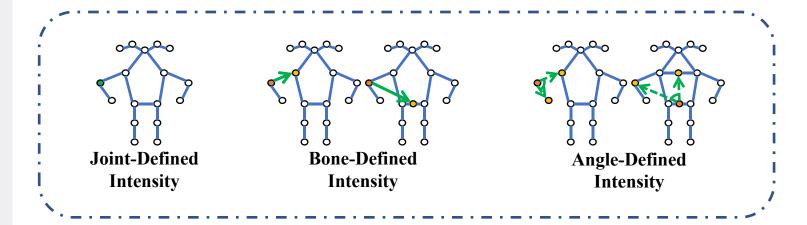
Bone-Defined

Angle-Defined

JBA-Defined

JBA-Defined

- The values of motion intensity calculated in three definitions need to be normalized because three definitions have different dimensions.
- Two kinds of methods to ensemble them:
- 1) Arithmetic average
- 2) Weighted average: utilize the normalized two-pathway accuracy we get from each definition as the weight of the corresponding definition



Datasets&Settings

Comparisons with SOTA Methods

Ablation Study

Datasets

1) FineGYM-99:

29K videos of 99 fine-grained gymnastic action classes

2) HMDB-51:

• 51 distinct action categories, each containing at least 101 clips for a total of 6766 video clips extracted from a wide range of sources

Settings

1) Pytorch framework, 4 NVIDIA GTX 3090 GPUs

2) For FineGYM-99:

• learning rate: 0.4, weight decay: 0.0003, learning rate adjustment: cosine annealing strategy, number of epochs: 80, model: trained from scratch, ResNet blocks setting is (4,6,3)

3) For HMDB-51:

• learning rate: 0.01, weight decay: 0.0001, learning rate adjustmentis: step strategy, number of epochs: 12, model: pretrained on Kinetics400, ResNet blocks setting is (3,4,6)

4) For both datasets:

• batch size: 16, optimizer: SGD, momentum: 0.9, the length of sampled clip: 48

The code has been shown at https://github.com/SarahQi666/Int-and-Int.

Datasets&Settings

Comparisons with SOTA Methods

Ablation Study

Comparisons with SOTA Methods

 Although these prior works all use at least videos as the input which contain much more information than our skeleton-based method, our results achieve superior performance and outperform the results of SOTA methods.

FineGYM-99

Method	Accuracy
TSN(RGB) [6] [19]	74.8%
TSN(RGB+Flow) [6] [19]	86.0%
TRN(RGB) [7] [19]	79.9%
TRN(RGB+Flow) [7] [19]	87.4%
ActionVLAD [9] [19]	69.5%
Int∬(Ours)	95.6%

For FineGYM-99, we report the Top-1 accuracy.
 For HMDB-51, we report average accuracy over three splits.

HMDB-51

Method	Accuracy
Two-Stream(fusion by SVM) [1]	59.4%
Two-Stream(fusion by averaging) [1]	58.0%
TDD [2]	63.2%
C3D [3] [8]	56.8%
TSN(RGB+Flow) [6]	68.5%
TSN(RGB+Flow+Warped Flow) [6]	69.4%
TLE(FC-Pooling) [8]	68.8%
ActionVLAD [9]	66.9%
LTC(RGB+Flow) [10]	64.8%
Int∬(Ours)	69.6%

Datasets&Settings

Comparisons with SOTA Methods

Ablation Study

Ablation Study

1) Single-pathway vs. Two-pathway

• The two-pathway architecture almost consistently outperforms the single-pathway architectures.

	HMDB-51											
Sampling	3 Splits			Split 1			Split 2			Split 3		
	Int∬	Intensity Pathway	Integrity Pathway	Int∬	Intensity Pathway	Integrity Pathway	Int∬	Intensity Pathway	Integrity Pathway	Int∬	Intensity Pathway	Integrity Pathway
Random	69.3%	63.7%	68.6%	70.2%	64.4%	69.2%	69.0%	63.0%	68.6%	68.6%	63.7%	68.0%
Joint-Defined	69.6%	63.9%	68.6%	70.5%	64.5%	69.2%	69.3%	63.5%	68.6%	69.0%	63.5%	68.0%
Bone-Defined	69.0%	64.2%	68.6%	69.1%	63.9%	69.2%	68.7%	63.7%	68.6%	69.2%	65.0%	68.0%
Angle-Defined	69.5%	63.4%	68.6%	69.3%	63.9%	69.2%	70.3%	62.6%	68.6%	68.8%	63.6%	68.0%
JBA(Arithmetic)	69.3%	63.6%	68.6%	68.6%	63.4%	69.2%	70.1%	63.2%	68.6%	69.2%	64.3%	68.0%
JBA(Weighted)	69.2%	63.9%	68.6%	69.6%	64.0%	69.2%	69.7%	64.5%	68.6%	68.2%	63.1%	68.0%

2) Randomly sampling vs. Intensity-dependent sampling

Randomly sampling: the central frame in the sampled —
clip is not the most intense frame of the video but a strame randomly selected.

Intensity-dependent sampling consistently outperforms = sampling from a random temporal window.

Sampling	FineGYM-99	HMDB-51					
Samping	Tilled TM-99	3 Splits	Split 1	Split 2	Split 3		
Random	95.3%	69.3%	70.2%	69.0%	68.6%		
Intensity	95.6%	69.6%	70.5%	70.3%	69.2%		

Datasets&Settings

Comparisons with SOTA Methods

Ablation Study

Ablation Study

3) Intensity defined through motion prior knowledge

- Various definitions place emphasis on different semantic information:
- a) The positions of joints carry the most direct and obvious information of motion.
- b) The lengths and directions of bones contain higher-order information, which have been proven to be effective in 2s-AGCN.
- c) Angles can maintain invariance against different human body sizes and discriminate actions sharing similar motion trajectories such as taking off glasses and taking off headphones.

Sampling	FineGYM-99	HMDB-51					
		3 Splits	Split 1	Split 2	Split 3		
Joint-Defined	95.0%	69.6%	70.5%	69.3%	69.0%		
Bone-Defined	95.2%	69.0%	69.1%	68.7%	69.2%		
Angle-Defined	95.5%	69.5%	69.3%	70.3%	68.8%		
JBA(Arithmetic)	95.5%	69.3%	68.6%	70.1%	69.2%		
JBA(Weighted)	95.6%	69.2%	69.6%	69.7%	68.2%		

- For the professional and flexible gymnastics, it is necessary to combine the information from three aspects because of the richness of fine-grained action information.
- For the actions in HMDB-51, which are generally simpler, the information from only one aspect is enough to capture the feature.

CONCLUSION

- In this work, we propose a two-pathway *Int&Int* network for skeleton-based action recognition. The final model achieves superior performance on both of the datasets.
- In the future:
- a) reduce the computational complexity of our network and further improve the performance
- b) conduct research on the applications such as security, human-computer interaction, physical training, medical rehabilitation, and entertainment

Thanks for your attention!

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