

Section 1: Introduction

- **Proposal:** a novel mid-level representation for action/activity recognition on RGB videos on the basis of improved dense trajectories (IDT) [1], fisher vectors (FV), and videodarwin (VD) [2].
- We model the evolution of features not only for the entire video, but also on its subparts (represented as nodes in a binary tree hierarchically grouping subsets of IDTs).
- For each node, we compute Node-VD and Branch-VD. These are later combined with with VD on the whole video trajectories (Root-VD) a to perform classification with SVM.
- Results: better performance than standard VD (i.e., global-VD) and defines the state-of-the-art on UCF-Sports [3] and Highfive [4] action recognition datasets.

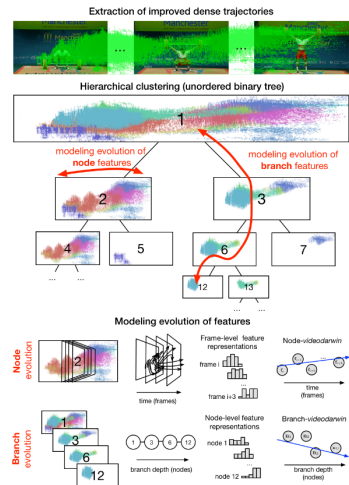


Fig. 1. The pipeline. Each leaf node is represented in a different color.

Section 2: Method

Binary tree of trajectory classification

- By recursively applying a *divisive spectral clustering algorithm* [5] on the set of trajectories D .
- For the clustering, we used primitive trajectory features $\bar{x}, \bar{y}, \bar{t}, \bar{v}_x, \bar{v}_y$.
- A tree node i containing the set of trajectories $D_i \subseteq D$ expands a temporal segment (t_i, t'_i) of the T -frame video, $0 \leq t_i < t'_i < T$.
- Let U_i and u_i be respectively the matrix of **per-frame FVs** and the **global FV** on D_i .

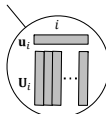


Fig. 2. i -th node representation: global FV for all IDTs assigned to the node's cluster, U_i , and matrix of per-frame FVs, u_i .

Videodarwin: in-a-nutshell

- VD applies any learning algorithm able to model frame ordering in a sequence. Our choice is to use a *linear regressor* we refer to as v .
- We compute VD in forward and reverse directions.
- Prior to VD, *time varying mean* is applied. Given $X \in \mathbb{R}^{\#(\text{features}) \times \#(\text{timesteps})}$, forward videodarwin (FW) is calculated as follows:

$$m_{\tau}^{FW} = \frac{1}{\tau} \sum_{k=1}^{\tau} X_{:,k}$$

$$V_{\tau}^{FW} = \frac{m_{\tau}^{FW}}{\|m_{\tau}^{FW}\|_1}, \forall \tau = 1, \dots, T$$

Note reverse VD simply re-defines m_{τ}^{FW} to calculate the varying mean backwards.

- The final VD representation, w , is then:

$$w^{FW} = v(V^{FW}, (1..T))$$

$$w^{RV} = v(V^{RV}, (1..T))$$

$$w = [(w^{FW})^T, (w^{RV})^T]^T$$

Mid-level representations

- **Node-VD** representation on node i , i.e. n_i , by taking $X = U_i$. In particular, **Root-VD** is just the special case $i = 1$.
- **Branch-VD** on node i requires its ancestors to be represented by their

global FV, u_i . We construct i -th node's branch as a matrix of per-node global FVs. That is:

$$B_i = [u_i, u_i / 2^i, u_i / 2^{2i}, \dots, u_i]$$

- Then, i -th node's branch representation, b_i , is computed taking $X = B_i$.

Darwintree kernel classification

- Each tree has an arbitrary number of nodes and each node is represented by the combination of Node- and Branch-VD:

$$s_i = [n_i; b_i], i > 1.$$
- We define the *Darwintree kernel* function k_{DT} between two trees (S, S') based on pairwise similarities of their nodes' representations:

$$k_{DT}(S, S') = \frac{1}{|S||S'|} \sum_{s_i \in S} \sum_{s_j \in S'} \phi(s_i, s_j),$$

$\forall i, j > 1$, where $\phi(\cdot, \cdot)$ can be any linear mapping function (e.g. dot product).

Since root node has no ancestors, we define a different kernel:

$$k_{root}(n_1, n'_1) = \phi(n_1, n'_1)$$

- Finally, a **linear SVM** performs classification using a linear combination of k_{DT} and k_{root} :

$$k_{final} = (1 - \alpha) k_{DT} + \alpha k_{root}.$$

Section 3: Results

- We validated our method in UCF-Sports [3] and Highfive [4] datasets.
- **Node-VD (N) and Branch-VD (B) against Darwintrees (DT):** DT provided superior performance than N or B on UCF-Sports. On Highfive, DT demonstrated its complementarity with Root-VD.

Method	UCF [3] (acc)	Highfive [4] (mAP)		
		F#1	F#2	TOTAL
N	85.11	76.55	70.41	73.48
B	80.85	76.25	72.53	74.39
DT (N+B)	91.49	76.04	70.37	73.21
Root+DT	91.49	79.24	72.32	75.78

Table 1. Node-VD (N) Branch-VD (B) versus Darwintrees (DT) and DT combined with root (Root+DT) at kernel level.

- We also compared to other **SOTA methods**.

Method	Accuracy (%)
Ours (Root+DT)	91.5
Karaman et al. (2014)	90.8
Ma et al. (2015)	89.4
Wang et al. (2013)	85.2
Ma et al. (2013)	81.7
Raptis et al. (2012)	79.3

Table 2. Results on UCF-Sports dataset.

Method	mAP
Ours (Root+DT)	75.8
Wang et al. (2015)	69.4
Karaman et al. (2014)	65.4
Ma et al. (2015)	64.4
Gaidon et al. (2014)	62.4
Ma et al. (2013)	36.9
Patron-Pérez et al. (2012)	42.4

Table 3. Results on UCF-Sports dataset.

Section 4: Conclusions

- A novel mid-level representation for action recognition on RGB videos.
- We modeled the evolution of features on both trajectory clusters and on the hierarchy defining those groupings.
- It is applicable to any local spatio-temporal feature representation.
- We demonstrated superior performance than other SOTA methods, especially for Highfive.

References

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