

Motivation

- Misalignment is caused by variations of viewpoints and poses, which commonly exist in person re-identification.
- Some local patches are more distinctive and reliable when matching two persons.
- Images of the same person captured from different camera views have some invariance property on their spatial distribution on saliency.

Contribution

- A probabilistic distribution of saliency is reliably estimated with our approach.
- We formulate person re-identification as a saliency matching problem.
- Saliency matching and patch matching are tightly integrated into a unified structural RankSVM learning framework.

Unsupervised Human Saliency Learning

Algorithm 1 Compute human saliency.

Input: image $\mathbf{x}^{A,u}$ and a reference image set $\mathcal{R} = \{\mathbf{x}^{B,v}, v = 1, \dots, N_r\}$

Output: saliency probability map $P(l_{m,n}^{A,u} = 1 | x_{m,n}^{A,u})$

- 1: for each patch $x_{m,n}^{A,u} \in X$ do
- 2: compute $X_{NN}(x_{m,n}^{A,u})$ with Eq. (1)
- 3: compute $score(x_{m,n}^{A,u})$ with Eq. (2)
- 4: compute $P(l_{m,n}^{A,u} = 1 | x_{m,n}^{A,u})$ with Eq. (3)
- 5: end for

Step 1: construct nearest-neighbor patch set

$$X_{NN}(x_{m,n}^{A,u}) = \{x | \operatorname{argmin}_{x_{i,j}^{B,v}} d(x_{m,n}^{A,u}, x_{i,j}^{B,v}), \quad (1)$$



$$x_{i,j}^{B,v} \in \mathcal{S}(x_{m,n}^{A,u}, \mathbf{x}^{B,v}), v = 1, \dots, N_r, \quad (2)$$

Step 2: compute k-NN distances as saliency score

$$score(x_{m,n}^{A,u}) = d_k(X_{NN}(x_{m,n}^{A,u})), \quad (2)$$

Step 3: convert to saliency probability

$$P(l_{m,n}^{A,u} = 1 | x_{m,n}^{A,u}) = 1 - \exp(-score(x_{m,n}^{A,u})^2 / \sigma_0^2), \quad (3)$$

Supervised Saliency Matching Framework

Matching based on Saliency:

$$f_z(\mathbf{x}^A, \mathbf{x}^B, \mathbf{l}^A, \mathbf{l}^B; \mathbf{p}, \mathbf{z}) = \sum_{p_i} \left\{ z_{p_i,1} l_{p_i}^A l_{p_i'}^B + z_{p_i,2} l_{p_i}^A (1 - l_{p_i'}^B) + z_{p_i,3} (1 - l_{p_i}^A) l_{p_i'}^B + z_{p_i,4} (1 - l_{p_i}^A) (1 - l_{p_i'}^B) \right\},$$

➤ **Hidden Saliency label:**

$$\mathbf{l}^A = \{l_{p_i}^A | l_{p_i}^A \in \{0, 1\}\} \quad \mathbf{l}^B = \{l_{p_i'}^B | l_{p_i'}^B \in \{0, 1\}\}$$

➤ **Matching score:** $\mathbf{z} = \{z_{p_i,k}\}_{k=1,2,3,4}$

$$z_{p_i,k} = \alpha_{p_i,k} \cdot s(x_{p_i}^A, x_{p_i'}^B) + \beta_{p_i,k}, \quad s(x_{p_i}^A, x_{p_i'}^B) = \exp\left(-\frac{d(x_{p_i}^A, x_{p_i'}^B)^2}{2\sigma_0^2}\right),$$

Marginalization:

$$f^*(\mathbf{x}^A, \mathbf{x}^B; \mathbf{p}, \mathbf{z}) = \sum_{\mathbf{l}^A, \mathbf{l}^B} f_z(\mathbf{x}^A, \mathbf{x}^B, \mathbf{l}^A, \mathbf{l}^B; \mathbf{p}, \mathbf{z}) p(\mathbf{l}^A, \mathbf{l}^B | \mathbf{x}^A, \mathbf{x}^B) = \sum_{p_i} \sum_{k=1}^4 \left[\alpha_{p_i,k} \cdot s(x_{p_i}^A, x_{p_i'}^B) + \beta_{p_i,k} \right] c_{p_i,k}(x_{p_i}^A, x_{p_i'}^B),$$

$$c_{p_i,k}(x_{p_i}^A, x_{p_i'}^B) = \begin{cases} P(l_{p_i}^A = 1 | x_{p_i}^A) P(l_{p_i'}^B = 1 | x_{p_i'}^B), & k = 1, \\ P(l_{p_i}^A = 1 | x_{p_i}^A) P(l_{p_i'}^B = 0 | x_{p_i'}^B), & k = 2, \\ P(l_{p_i}^A = 0 | x_{p_i}^A) P(l_{p_i'}^B = 1 | x_{p_i'}^B), & k = 3, \\ P(l_{p_i}^A = 0 | x_{p_i}^A) P(l_{p_i'}^B = 0 | x_{p_i'}^B), & k = 4. \end{cases}$$

Probabilistic saliency matching

Final Formulation:

$$f^*(\mathbf{x}^A, \mathbf{x}^B; \mathbf{p}, \mathbf{z}) = \mathbf{w}^T \Phi(\mathbf{x}^A, \mathbf{x}^B; \mathbf{p}) = \sum_{p_i} w_{p_i}^T \phi(x_{p_i}^A, x_{p_i'}^B),$$

➤ **Weighting parameters:** $w_{p_i} = [\{\alpha_{p_i,k}\}_{k=1,2,3,4}, \{\beta_{p_i,k}\}_{k=1,2,3,4}]$.

➤ **Matching feature:** $\phi(x_{p_i}^A, x_{p_i'}^B)$

$$\phi(x_{p_i}^A, x_{p_i'}^B) = \begin{bmatrix} s(x_{p_i}^A, x_{p_i'}^B) P(l_{p_i}^A = 1 | x_{p_i}^A) P(l_{p_i'}^B = 1 | x_{p_i'}^B) \\ s(x_{p_i}^A, x_{p_i'}^B) P(l_{p_i}^A = 1 | x_{p_i}^A) P(l_{p_i'}^B = 0 | x_{p_i'}^B) \\ s(x_{p_i}^A, x_{p_i'}^B) P(l_{p_i}^A = 0 | x_{p_i}^A) P(l_{p_i'}^B = 1 | x_{p_i'}^B) \\ s(x_{p_i}^A, x_{p_i'}^B) P(l_{p_i}^A = 0 | x_{p_i}^A) P(l_{p_i'}^B = 0 | x_{p_i'}^B) \\ P(l_{p_i}^A = 1 | x_{p_i}^A) P(l_{p_i'}^B = 1 | x_{p_i'}^B) \\ P(l_{p_i}^A = 1 | x_{p_i}^A) P(l_{p_i'}^B = 0 | x_{p_i'}^B) \\ P(l_{p_i}^A = 0 | x_{p_i}^A) P(l_{p_i'}^B = 1 | x_{p_i'}^B) \\ P(l_{p_i}^A = 0 | x_{p_i}^A) P(l_{p_i'}^B = 0 | x_{p_i'}^B) \end{bmatrix}.$$

Ranking by Partial Order

➤ **Task – finding a good ranking:**

$$\mathbf{y}_*^{A,u} = \operatorname{argmax}_{\mathbf{y}^{A,u} \in \mathcal{Y}^{A,u}} \mathbf{w}^T \Psi_{po}(\mathbf{x}^{A,u}, \mathbf{y}^{A,u}; \{\mathbf{x}^{B,v}\}_{v=1}^V, \{\mathbf{p}^{u,v}\}_{v=1}^V),$$

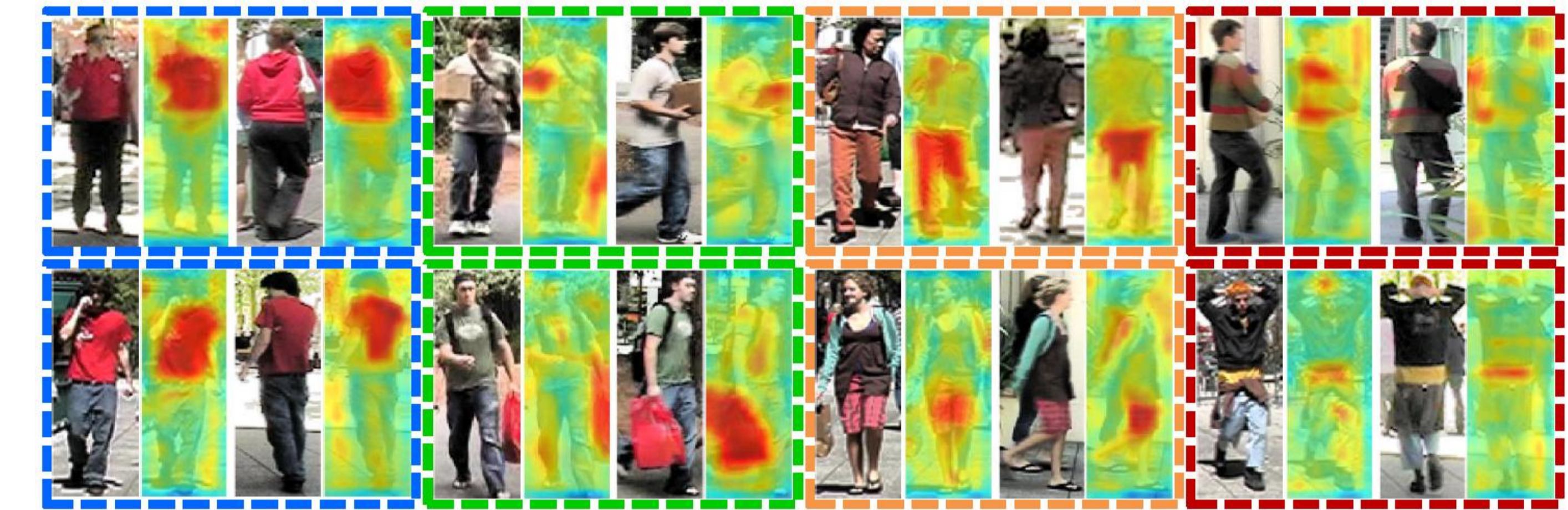
Partial order feature for structural RankSVM training:

$$\Psi_{po}(\mathbf{x}^{A,u}, \mathbf{y}^{A,u}; \{\mathbf{x}^{B,v}\}_{v=1}^V, \{\mathbf{p}^{u,v}\}_{v=1}^V) = \sum_{\mathbf{x}^{B,v} \in S_{\mathbf{x}^{A,u}}^+} \sum_{\mathbf{x}^{B,v'} \in S_{\mathbf{x}^{A,u}}^-} y_{v,v'}^{A,u} \frac{\Phi(\mathbf{x}^{A,u}, \mathbf{x}^{B,v}; \mathbf{p}^{u,v}) - \Phi(\mathbf{x}^{A,u}, \mathbf{x}^{B,v'}; \mathbf{p}^{u,v'})}{|S_{\mathbf{x}^{A,u}}^+| \cdot |S_{\mathbf{x}^{A,u}}^-|},$$

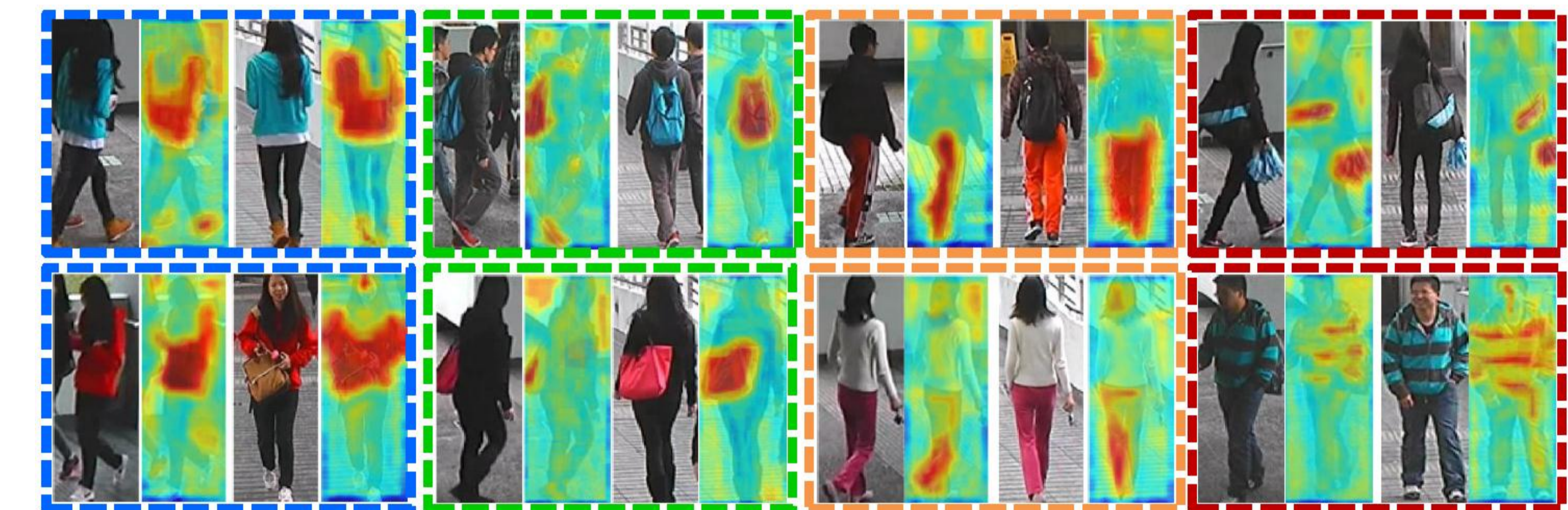
➤ **Solution – sorting gallery by $\{\mathbf{w}^T \Phi(\mathbf{x}^{A,u}, \mathbf{x}^{B,v}; \mathbf{p}^{u,v})\}_v$ in descending order.**

Experimental Results

Learned Human Saliency

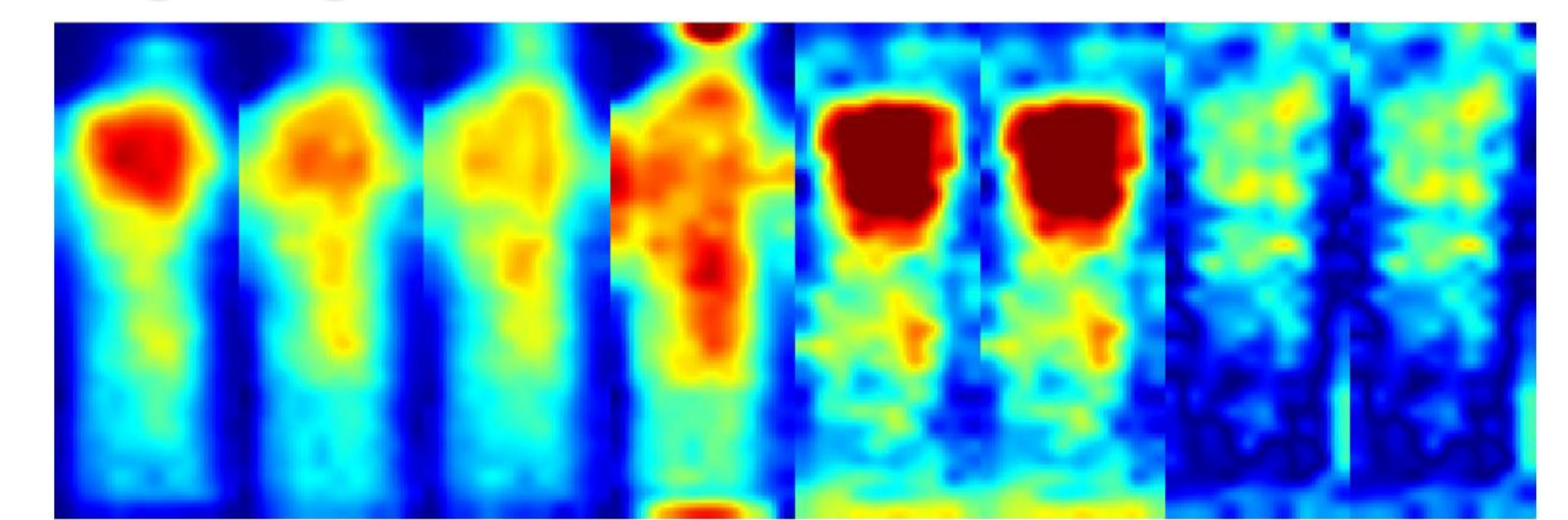


(a) VIPeR dataset



(b) CUHK Campus dataset

Learned Weighting Parameters:



$\alpha_{p_i,1} \quad \alpha_{p_i,2} \quad \alpha_{p_i,3} \quad \alpha_{p_i,4} \quad \beta_{p_i,1} \quad \beta_{p_i,2} \quad \beta_{p_i,3} \quad \beta_{p_i,4}$

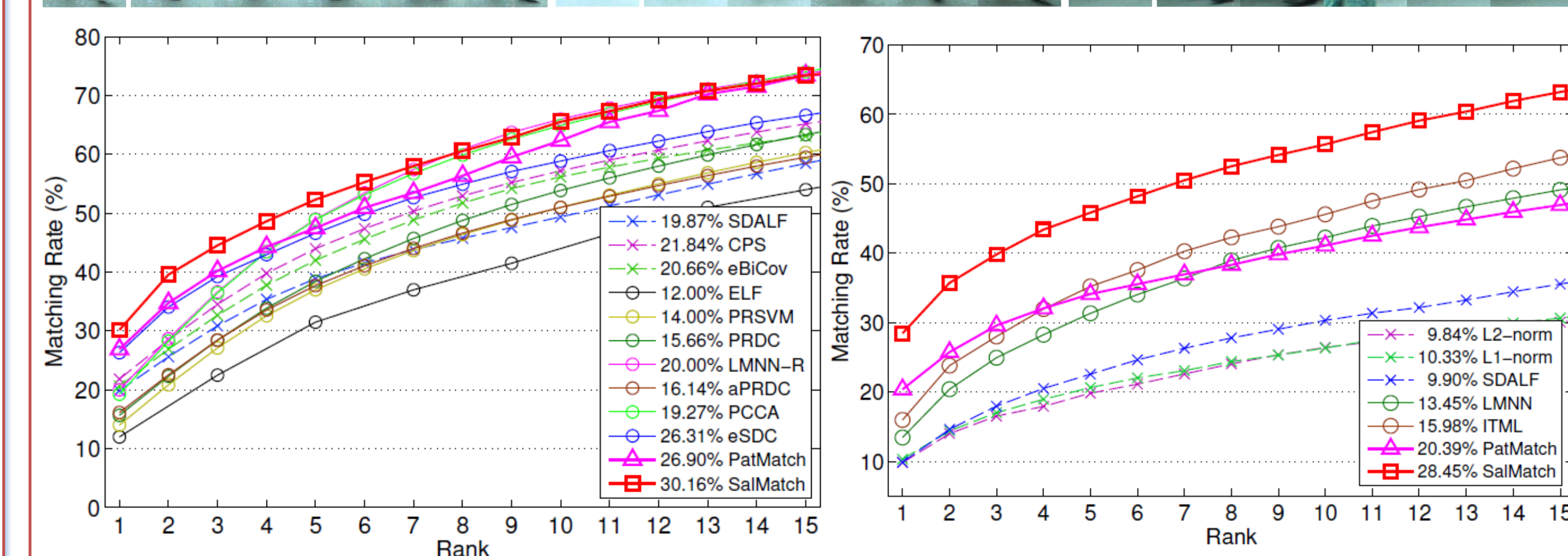
$\{\alpha_{p_i,k}\}_{k=1,2,3,4}$ correspond to the first four terms of matching features based on saliency matching with visual similarity, and $\{\beta_{p_i,k}\}_{k=1,2,3,4}$ correspond to the last four terms only depending on saliency matching.

Comparison with State-of-the-Arts:

VIPeR dataset



CUHK Campus dataset



(a) VIPeR dataset

(b) CUHK Campus dataset