

# **Group-aware Label Transfer for Domain Adaptive Person Re-identification**

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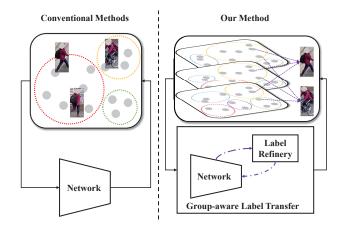
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# **Abstract**

Unsupervised Domain Adaptive (UDA) person reidentification (ReID) aims at adapting the model trained on a labeled source-domain dataset to a target-domain dataset without any further annotations. Most successful UDA-ReID approaches combine clustering-based pseudo-label prediction with representation learning and perform the two steps in an alternating fashion. However, offline interaction between these two steps may allow noisy pseudo labels to substantially hinder the capability of the model. In this paper, we propose a Group-aware Label Transfer (GLT) algorithm, which enables the online interaction and mutual promotion of pseudo-label prediction and representation learning. Specifically, a label transfer algorithm simultaneously uses pseudo labels to train the data while refining the pseudo labels as an online clustering algorithm. It treats the online label refinery problem as an optimal transport problem, which explores the minimum cost for assigning M samples to N pseudo labels. More importantly, we introduce a group-aware strategy to assign implicit attribute group IDs to samples. The combination of the online label refining algorithm and the group-aware strategy can better correct the noisy pseudo label in an online fashion and narrow down the search space of the target identity. The effectiveness of the proposed GLT is demonstrated by the experimental results (Rank-1 accuracy) for Market1501→DukeMTMC (82.0%) and DukeMTMC→Market1501 (92.2%), remarkably closing the gap between unsupervised and supervised performance on person re-identification.

### 1. Introduction

Person re-identification (ReID) [27, 50, 19] is the important task of matching person images captured from non-overlapping camera networks, which is widely used in prac-



**Figure 1.** Illustration of conventional methods and our group-aware label transfer method. In our method, each instance is assigned to multiple prototypes with different granularity for generating multi-group pseudo labels, and then the noisy multi-group pseudo labels are online refined. With guidance of refined multigroup pseudo labels, our method can learn an embedding space that encodes the semantic multi-granularity structure of data.

tical applications such as automatic surveillance, content-based retrieval, and behavior analysis [7, 23, 52, 48, 44, 24, 30, 12, 28, 17, 29]. It has been proved that existing approaches can achieve remarkable performance when the training and testing data are collected from the same application scenario but often fail to generalize well to other scenarios due to the inevitable domain gaps. Therefore, it is necessary for both academia and industry to study the Unsupervised Domain Adaptive (UDA) person re-identification problem.

Existing UDA ReID approaches [14, 26, 35, 45, 55] typically include three steps: feature pre-training with labeled source domain data, clustering-based pseudo-label prediction for the target domain data, and feature representation learning/fine-tuning with the pseudo-labels. The last two steps are usually iteratively conducted to strengthen or promote each other. However, the first problem is that the pseudo-labels assigned through clustering usually contain

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<sup>&</sup>lt;sup>1</sup>Full codes are available in https://github.com/JDAI-CV/fast-reid and https://github.com/zkcys001/UDAStrongBaseline.

incorrect labels due to the divergence/domain gap between the source and target data, and the imperfect nature of the clustering algorithm. Such noisy labels may mislead the feature learning and harm the domain adaptation performance. Although the label refining objective in clustering is tractable, it is an offline time-consuming scheme as it requires a pass over the entire dataset. Therefore, online refining those incorrect samples when training can help model learn more robust and accurate representation.

Another problem is that the target domain lacks ID information, so it is difficult to cluster the person images according to human identity. However, in the real world, each person has their own characteristics in his or her appearance. There may be common appearances shared by a group of people but they are not the same identity (e.g. two men with the same red coats and similar black pants as shown in Fig. 1). Therefore, group-based description [21] that involves common characteristics in a pseudo group, can be useful to narrow down the set of candidates and beneficial to identify the exact persons [47, 3, 2]. This group-aware strategy can cluster a person into multi-group clustering prototypes, and is able to efficiently embed a significant number of people and briefly describe an unknown person. Inspired by this, as shown in Fig. 1, combining the online label refining algorithm with the group-aware strategy may be beneficial to the success of domain adaptation.

In this paper, we propose a Group-aware Label Transfer (GLT) algorithm that facilitates the online interaction and mutual promotion of the pseudo labels prediction and feature learning. Specifically, our method simultaneously uses pseudo labels to train the data while online refining the pseudo labels via the label transfer algorithm. This label transfer algorithm regards the resulting label refining problem as optimal transport, which explores the minimum cost for assigning M samples to N pseudo labels. This problem can therefore be solved by the Sinkhorn-Knopp algorithm [6] of linear programming in polynomial time. Meanwhile, the label transfer algorithm is scalable, which can be trained with several batches or the whole dataset and can scale to unlimited amounts of data. More importantly, we introduce a group-aware strategy to assign implicit attribute group IDs to samples. Explicit grouping requires manual annotation. Therefore, we adopt a group-aware clustering algorithm to generate multi-group pseudo labels. By adopting the concept of grouping, the ReID network can reduce the search space and flexibly embed a significant number of identities into an embedding feature. As shown in Fig. 1, we combine the online label refining algorithm with the multigroup pseudo labels. The combination of the online label refining algorithm and the group-aware strategy can better correct the noisy pseudo label online and narrow down the search space of the target identity and predict more accurate pseudo labels. In addition, we design a target instance memory bank combined with weighted contrastive loss [36] to make the model more powerful for features representation. The proposed GLT framework achieves state-of-the-art performance on Market-to-Duke, Duke-to-Market, Market-to-MSMT, and Duke-to-MSMT with unsupervised domain adaptation, and in particular the *i.e.* Duke-to-Market performances (92.2 Rank-1 and 79.5 mAP) are almost comparable with the supervised learning performances (94.1 Rank-1 and 85.7 mAP). Our method significantly closes gap between unsupervised and supervised performance on person re-identification, i.e., 92.2 vs. 94.1 top-1 accuracy in DukeMTMC to Market1501 transfer.

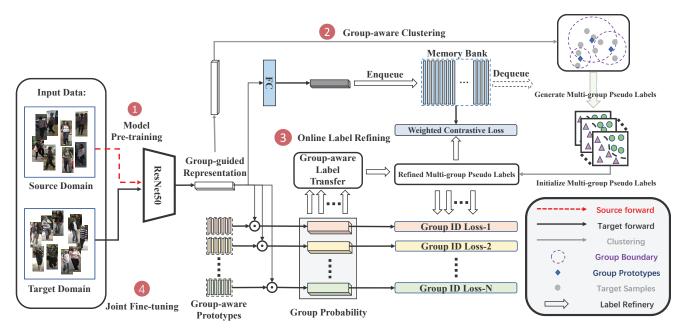
The main contributions of this paper can be summarized in four aspects:

- We make the first attempt towards integrating clustering and feature learning in a unified framework through the label transfer method for UDA-ReID. It can online refine the predicted pseudo labels to improve the feature representation ability of the model on the target domain.
- We propose a group-aware feature learning strategy based on label transfer to refine multi-group pseudo labels, which provides good latent pseudo-label groups for improving the quality of representation learning.
- The GLT framework achieves significant performance improvements compared to state-of-the-art approaches on Market—Duke, Duke—Market, Market—MSMT, Duke—MSMT ReID tasks. Even for the supervised learning methods, our algorithm is remarkably closing the gap.

# 2. Related Work

Clustering-based Methods in general generate hard or soft pseudo labels based on clustering results and then fine-tune/train the models based on the pseudo labels [11, 51, 42, 14, 43, 55, 13, 20, 4, 8, 49], which are widely used due to their superior performance. PUL [11] iteratively obtains the hard clustering labels and trains the model in a self-training manner. SSG [42] exploits the potential similarity of the global and local body parts, respectively, to build multiple independent clusters. PAST [51] develop a self-training method with progressive augmentation framework to promote the model performance progressively.

Pseudo label noises caused by unsupervised clustering are always an obstacle to UDA Re-ID. Such noisy labels would mislead the feature learning and impede the achievement of high performance. Recently, some methods introduce mutual learning among two or three collaborative networks to mutually exploit the refined soft pseudo labels of the peer networks as supervision [14, 46]. To suppress the noises in the pseudo labels, NRMT [53] maintains two



**Figure 2.** Illustration of our group-aware label transfer framework (GLT), where the group-aware label transfer module and the ReID model are alternately optimized to improve each other towards the final objective of accurate person re-ID. In the model pre-training stage, we pre-train the network on source domain labeled data. Then, in the group-aware clustering stage, we perform clustering algorithm on the unlabeled target data using the multi-group strategy and assign multi-group pseudo labels based on the clustering results. In the label online refining stage, the label transfer algorithm treats the resulting label refining problem as an optimal transport problem, and refine the multi-group pseudo labels by Sinkhorn-Knopp algorithm [6] of linear programming. In the joint fine-tuning stage, we use the refined multi-group pseudo labels to train the networks. Stage 3 and Stage 4 are performed alternatively in the online scheme.

networks during training to perform collaborative clustering and mutual instance selection, which reduces the fitting to noisy instances by using mutual supervision and the reliable instance selection. These approaches need mutual learning of two or more networks and are somewhat complicated. Besides, the selection of reliable instances in NRMT [53] is a hard selection, which requires a careful determination of threshold parameters and may lose the chance of exploiting useful information of the abandoned samples.

**Domain translation based methods** [9, 40] consider to translate target images by the source images. Then they use these translated source-to-target images and their corresponding ground-truth identities to fine-tune the target-domain model. PTGAN [40] utilizes the color consistency to impose pixel-level constraints during the domain translation. SPGAN [9] further minimizes the feature-level similarities between translated images and the original ones. This kind of approach does not apply to real-world scenarios, because the GAN-based method is hardly used to train and may introduce extra computing cost.

**Memory bank based methods** have been widely used for unsupervised representation learning which facilitates the introduction of contrastive loss [15] for the general tasks. He *et al.* [15] leverage the memory bank to better train the model to exploit the similarity between a sample and the

instances in the global memory bank. Wang et al. [39] propose using the memory bank to facilitate hard negative instance mining across batches. ECN [55] leverages memory bank to enforce exemplar-invariance, camera-invariance, and neighborhood-invariance over global training data (instead of the local batch) for UDA person ReID. We introduce contrastive loss to the target instance memory bank to enable the joint optimization of positive pairs and negative pairs for a query/anchor sample over all the samples in the memory bank, which serves as our strong baseline.

#### 3. Group-aware Label Transfer

We first introduce the overall architecture, then derive our group-aware label transfer framework and finally introduce the memory bank with weighted contrastive loss.

# 3.1. Clustering-based Method Revisit

**Formulation.** Unsupervised domain adaptive person reidentification aims at adapting the model trained on a source domain dataset with annotations to another unlabeled target domain dataset. Formally, given the source domain data  $\mathbb{D}_s = \left\{ \left. (\boldsymbol{x}_i^s, \boldsymbol{y}_i^s) \right|_{i=1}^{N_s} \right\} \text{ with the manual annotation, we can access target domain images } \mathbb{D}_t = \left. \left\{ \left. \boldsymbol{x}_i \right|_{i=1}^{N_t} \right\} \right. \text{ without any }$ 

ground-truth labels.  $N_s$  and  $N_t$  denote the number of the labeled source data and the unlabeled target data, respectively.

**Overview.** Deep-cluster framework is the general pipeline due to the conciseness and effectiveness of this framework. To be specific, a ReID model  $F(\cdot|\boldsymbol{\theta})$  firstly is pre-trained via a supervised task on source domain images, where  $\boldsymbol{\theta}$  denotes the parameters of network. Then we adopt this well-pretrained model to extract the features of target domain images which denote as  $\{f_i = F(x_i|\boldsymbol{\theta})\}|_{i=1}^{N_t} \in \mathbb{R}^D$ . After that, the target domain images are grouped into K classes by clustering algorithm. Let  $\{\tilde{\boldsymbol{y}}_i\}|_i^{N_t} \in \{1,\dots,K\}$  denote the pseudo labels generated by target domain images  $\{x_i\}|_i^{N_t}$ , which contain the noisy label due to the imperfect nature of clustering algorithm.

To calculate a identification loss, a classification head  $FC_k: \mathbb{R}^D \to \mathbb{R}^K$  is adopted to convert the embedding feature to a score vector, in the form of  $p(\tilde{y}_i|x_i) = \operatorname{softmax}(FC(F(x_i|\theta)))$ . The network parameters  $\theta$  and a learnable target-domain classifier  $FC_k$  are then optimized with respect to a cross-entropy loss  $\mathcal{L}_{id}(\theta)$  and a triplet loss  $\mathcal{L}_{tri}(\theta)$  in the form of,

$$\mathcal{L}_{id}(\boldsymbol{\theta}) = -\frac{1}{m_t} \sum_{i=1}^{m_t} \log p\left(\tilde{y}_i | \boldsymbol{x}_i\right),$$

$$\mathcal{L}_{tri}(\boldsymbol{\theta}) = \frac{1}{m_t} \sum_{i=1}^{m_t} \max\left(0, s_i^n - s_i^p + m\right),$$
(1)

where  $m_t$  is the mini-batch size of target data,  $s_i^p$  denotes the hardest positive cosine similarity,  $s_i^n$  denotes the hardest negative cosine similarity, scripts i, p and j, n indicate the indices of hardest positive and negative feature in each mini-batch for the sample  $x_i$ , and m=0 denotes the triplet distance margin.

Such two steps, pseudo label generation by clustering and feature learning with pseudo labels, are alternated until the training converges. However, because of the domain gap, the pseudo-labels are not always reliable and there are noisy labels. This would mislead the feature representation learning towards learning noisy information. So, we need to adopt an online clustering strategy to refine those samples with wrong pseudo labels. As illustrated in Fig. 2, the framework is alternated to gradually train the ReID model on the target domain with the guidance of pseudo labels.

# 3.2. Online Clustering by Group-aware Label Transfer

Each image  $x_i$  is transformed into  $f_i$  by a ReID model. We then compute a group probability  $p_i$  from this feature by mapping  $f_i$  to a set of K trainable prototype vectors,

 $C = \{c_1, c_2, \dots c_K\}$ . We denote the matrix by C whose columns are the  $\{c_1, c_2, \dots c_K\}$ . Then, the pseudo labels  $Q \in [0, 1]^{K \times N}$  are adopted to supervise the group probability P, and we use Label Transfer to online update the pseudo labels as training goes on. We now first introduce how to update the prototypes online.

**Prototype Prediction Problem.** Our prototypes can be implemented by a non-parametric prototypical classifier or a parametric linear classifier. The optimization can be viewed to optimize the cluster-assignment probability through the cross-entropy loss, where the prototypes  $\boldsymbol{C}$  represent the trainable weights for a linear classifier. With K-means clustering, the linear classifier has a fixed set of weights as the mean vectors for the representations in each cluster.

The prototypes are learned by the non-parametric or a parametric way noted above. Here, we will introduce the prototype prediction problem. Firstly, we rewrite the crossentropy loss of eq. (1) by encoding the labels as posterior distributions  $q\left(\tilde{y}_{i}|\boldsymbol{x}_{i}\right)\in\boldsymbol{Q}$ , which is generated by clustering algorithm:

$$\mathcal{L}_{g}(p, q; \boldsymbol{\theta}) = -\frac{1}{N} \sum_{N} q\left(\tilde{y}_{i} | \boldsymbol{x}_{i}\right) \log p\left(\tilde{y}_{i} | \boldsymbol{x}_{i}\right),$$
where  $p\left(\tilde{y}_{i} | \boldsymbol{x}_{i}\right) = \frac{exp\left(\frac{1}{\tau} \boldsymbol{f}_{i} \boldsymbol{c}_{k}\right)}{\sum_{k'} exp\left(\frac{1}{\tau} \boldsymbol{f}_{i} \boldsymbol{c}_{k'}\right)},$ 
(2)

where  $\tau$  is a temperature parameter. If we set the posterior distributions  $q\left(\tilde{y}_i|\mathbf{x}_i\right) = \delta\left(\tilde{y}_i - y_i\right)$  to be deterministic, this eq. (2) is similar to the general cross-entropy loss. We can use a non-parametric prototypical classifier performs or a parametric linear classifier to update the prototypes. This loss function is jointly minimized respect to the prototypes C and the image encoder parameters  $\theta$ .

Online Refining Pseudo Labels via Label Transfer. the Clustering-based UDA methods, inevitable noisy labels caused by the clustering procedure bring a negative effect on network training. To address this issue, we propose a label transfer method to correct the noisy pseudo labels in an online training scheme by combining the representation learning and clustering algorithm. The general pipeline alternately tackles the optimization of q and representational learning p. However, separate optimizations cannot well generate accurate pseudo labels in representational learning because of restraining the interaction of these two steps. In order to make our method online, we try to utilize the prototypes and group probability to refine noisy pseudo labels. We compute refinery pseudo labels using the prototypes Csuch that all the examples in a batch are equally partitioned by the prototypes. This equipartition constraint ensures that the refinery pseudo labels for different images in several batches or a whole dataset are distinct, which preventing the trivial solution where every image has the same pseudo label:

$$\min_{p,q} \mathcal{L}_{g}(p,q;\boldsymbol{\theta}) \,\forall \tilde{y}_{i} : q\left(\tilde{y}_{i}|\boldsymbol{x}_{i}\right) \in \{0,1\}$$

$$\sum_{i=1}^{N} q\left(\tilde{y}_{i} = k|\boldsymbol{x}_{i}\right) = \frac{N}{k},$$
(3)

where each sample  $x_i$  is assigned to exactly one label and the N data points are split among the K classes for alleviating a trivial solution. Inspired by [1], they enforce an equal partition by constraining the matrix  $\mathbf{Q}$  to belong to the transportable prototype. We also propose to adapt their solution to work onthe whole dataset by restricting the transportable prototype.

The objective in eq. (3) is combinatorial in q and thus is very difficult to optimize by an end-to-end manner. However, the optimal transport algorithm [6] is able to solve this issue. Let  $P_{\tilde{y}_i,i} = p\left(\tilde{y}_i|\boldsymbol{x}_i\right)\frac{1}{N}$  be the  $(i,\tilde{y}_i)$  element of  $K\times N$  matrix as the joint probabilities estimated by the model, which represents the group probability that the i-th sample is predicted to become the  $\tilde{y}_i$ -th pseudo-label. Similarly,  $Q_{y_i,i} = q\left(\tilde{y}_i|\boldsymbol{x}_i\right)\frac{1}{N}$  is  $K\times N$  matrix as the pseudo label matrix. Using the notation of [6], we relax matrix  $\boldsymbol{Q}$  to be an element of the transportable prototype:

$$H(\boldsymbol{w}, \boldsymbol{c}) := \{ \boldsymbol{Q} \in \mathbb{R}_{+}^{K \times N} | \boldsymbol{Q} \cdot \mathbf{1} = \boldsymbol{w}, \boldsymbol{Q}^{\top} \cdot \mathbf{1} = \boldsymbol{c} \},$$

$$\boldsymbol{c} = \frac{1}{N} \cdot \mathbf{1}, \boldsymbol{w} = \frac{1}{N} \cdot \mathbf{1},$$
(4)

where 1 are vectors of all ones of the appropriate dimensions, so that c and w are the marginal projections of matrix Q onto its rows and columns, respectively. In our case, we require Q to be a matrix of conditional probability distributions that is similar to the number distribution of source domain.

In practice, the computational complexity is enormous, including millions of data points and thousands of classes. Traditional algorithms for solving this problem are difficult to be applied. Thus, we adopt the Sinkhorn-Knopp algorithm [6] to address this issue. This amounts to introducing a regularization term:

$$\min_{\boldsymbol{Q} \in H(\boldsymbol{w}, \boldsymbol{c})} \langle \boldsymbol{Q}, -\log \boldsymbol{P} \rangle + \frac{1}{\lambda} \operatorname{H}(\boldsymbol{Q}), \qquad (5)$$

where  $\langle . \rangle$  is the Frobenius dot-product between two matrices,  $\mathbf{R}$  denotes to the number distribution of source domain,  $\langle \mathbf{Q}, -\log \mathbf{P} \rangle = -\frac{1}{N} \sum_{i=1}^{N} \sum_{\tilde{y}_i=1}^{K} q\left(\tilde{y}_i|\mathbf{x}_i\right) \log p\left(\tilde{y}_i|\mathbf{x}_i\right)$ , log is applied element-wise,  $\mathbf{H}$  is the entropy function,  $\mathbf{H}(Q) = -\sum_{q \in \mathbf{Q}} q \log q$  and  $\lambda$  is a parameter that controls the smoothness of the mapping. We find that using a low  $\lambda$  can have a strong entropy regularization, which generally leads to a trivial solution where all samples collapse into an

# Algorithm 1 Group-aware Label Transfer

```
Require: Source domain data \mathbb{D}_s = \left\{ \left. (\boldsymbol{x}_i^s, \boldsymbol{y}_i^s) \right|_{i=1}^{N_s} \right\}, target domain images
       \mathbb{D}_{t} = \left\{ \left. \boldsymbol{x}_{i} \right|_{i=1}^{N_{t}} \right\}, \text{multi-group } k \text{ values } \left\{ k_{1}, \dots, k_{M} \right\};
 1: Pre-training source-domain encoder F(\cdot|\theta) by minimizing Eq. (1) on \mathbb{D}_s;
 2: Generating multi-group pseudo labels \{m{Q}_{k_m}\}|_{m=1}^M by clustering the target
       domain features \{F(\boldsymbol{x}_i|\boldsymbol{\theta})\}|_{i=1}^{N_t}.;
 3: for i in [1, iterations] do
             Updating parameters \theta by the gradient descent of objective function equation (11) under the guidance of multi-group pseudo labels \{Q_{k_m}\}_{m=1}^{|M|};
             for k_m in \{k_1, \ldots, k_M\} do
                    Extracting the multi-group probability matrix oldsymbol{P}_{k_m} of target images \mathbb{D}_t
6:
                    by the current model F(\cdot|\boldsymbol{\theta});
7:
                   \begin{aligned} & \text{while } ||\alpha_{j-1} - \alpha_{j}||_{1} < 0.1 \text{ do} \\ & \alpha_{j} \leftarrow \left[P_{k_{m}}^{\lambda} \beta_{j-1}\right]^{-1}, \beta_{j} \leftarrow \left[\alpha_{j-1}^{\top} P_{k_{m}}^{\lambda}\right]^{-1}; \end{aligned}
8:
9:
                                 =\operatorname{diag}(oldsymbol{lpha})oldsymbol{P}_{k_m}^{\lambda}\operatorname{diag}(oldsymbol{eta}) # Refining noisy multi-group
10:
11:
             Refining the multi-group pseudo labels Q_{k_m} = Q_{k_m}';
12: end for
```

unique representation and are all assigned uniformely to all prototypes. Hence, in practice we keep a high  $\lambda$ . According to the Sinkhorn-Knopp algorithm [1], this minimizer of regularization term of eq. (5) can be written as:

$$Q = \operatorname{diag}(\alpha) P^{\lambda} \operatorname{diag}(\beta) = \operatorname{diag}(\alpha) (C^{T} F)^{\lambda} \operatorname{diag}(\beta),$$
(6)

where  $F = [f_1, \ldots, f_N]$  denotes to the feature vectors of images,  $C = [c_1, \ldots, c_K]$  denotes to the prototypes of pseudo labels,  $\alpha$  and  $\beta$  are two renormalization vectors of scaling coefficients, the resulting matrix Q is also a probability matrix that represent the refining pseudo labels (see [6] for a derivation). By the iterative Sinkhorn-Knopp algorithm [1], the vectors  $\alpha$  and  $\beta$  can be obtained via a simple matrix scaling iteration:

$$\forall y : \alpha_y \leftarrow \left[ \mathbf{P}^{\lambda} \beta \right]_{y}^{-1} \quad \forall i : \beta_i \leftarrow \left[ \alpha^{\top} \mathbf{P}^{\lambda} \right]_{i}^{-1}, \quad (7)$$

where  $\alpha_0 = c$  and  $\beta_0 = w$ . When  $\lambda$  is very large, optimizing eq. (7) is mostly equivalent to optimizing eq. (6). However, even for intermediate values of lambda, the results still tend to approximate the optimal solution [6]. In the original optimal transport problem,  $\lambda$  adjust the balance between the convergence speed and optimum approximation. In our case, we only regard the  $\lambda$  as the hype-parameter, because we are more interested in obtaining a better performance in the final clustering and representation learning results, rather than solving the optimal transport problem.

After using a group probability matrix P to update the pseudo label matrix Q, we use the current Q as the pseudo label to train the ReID model for optimizing the group probability P. Such two steps, pseudo label prediction and feature learning with pseudo labels are alternated until the training converges.

**Group-aware Label Transfer.** The actual identification number k of the target domain is unknown, and it is hard to use the one value of k to generate accurate pseudo labels. Therefore, group-based description [21] that involves common characteristics in the pseudo groups, can be useful to narrow down the set of candidates, which is beneficial to identify the exact unlabeled persons. This groupaware strategy can cluster a person into multi-group clustering prototypes, which is able to efficiently embed a significant number of people and briefly describe an unknown person. Then, we introduce the group-aware strategy to assign implicit attribute group IDs to samples.

In our formulation, we firstly adopt the multi-k setting  $\{k_1,\ldots,k_M\}$  to generate the multi-group pseudo label via multi-k K-means. Then, we use them to train the ReID model:

$$\mathcal{L}_{G} = -\frac{1}{N} \sum_{m=1}^{M} \sum_{N} q_{k_{m}} \left( \tilde{y}_{i} | \boldsymbol{x}_{i} \right) \log p_{k_{m}} \left( \tilde{y}_{i} | \boldsymbol{x}_{i} \right), \quad (8)$$

Based on the multi-group pseudo labels, the group-aware label transfer module adopts the multi-group optimization strategy to refine these pseudo labels. The different groups of pseudo labels have different classification heads, while sharing the parameters of feature extractor  $\theta$  among them. In formulation, we optimize the multi-group objective functions of the eq. (9).

$$\begin{split} \sum_{m=1}^{M} \min_{\boldsymbol{Q}_{k_m} \in H_{k_m}} (\langle \boldsymbol{Q}_{k_m}, -\log \boldsymbol{P}_{k_m} \rangle + \frac{1}{\lambda} \mathrm{H}(\boldsymbol{Q}_{k_m})), \\ H_{k_m} := \left\{ \boldsymbol{Q}_{k_m} \in \mathbb{R}_{+}^{K_m \times N} | \boldsymbol{Q}_{k_m} \cdot \mathbf{1} = \boldsymbol{r}, \boldsymbol{Q}_{k_m}^{\top} \cdot \mathbf{1} = \boldsymbol{c} \right\}, \end{split}$$

$$\boldsymbol{c} = \frac{1}{N} \cdot \mathbf{1}, \boldsymbol{r} = \frac{1}{N} \cdot \mathbf{1}, \end{split}$$

where  $\{P_{k_m} = C_{k_m}^T \cdot F\}_{m=1}^M$  denotes to the multi-group probability and  $\{C_{k_m}\}_{m=1}^M$  denotes to the multi-group prototypes. Each update involves M matrix-vector multiplication with complexity  $\mathcal{O}(MNK)$ , so it is relatively quick.

#### 3.3. Target Instance Memory Bank

We first describe our target instance memory bank module, with an updating mechanism. Then we show that a weighting scheme focusing on informative pairs is beneficial for contrastive loss with multiple positive samples.

**Updating mechanism.** We adopt the memory bank [16] to remember the previous feature embeddings. We maintain the memory bank as a queue of data samples while minibatches of samples arrive. This allows us to reuse feature embeddings from the immediately preceding mini-batch. The size of our memory banks can be much larger than the typical mini-batch size and can be set flexibly and independently to hyperparameters. The samples in the memory bank are progressively replaced. In addition, the current

mini-batch are enqueued into the dictionary and the oldest mini-batch in the queue are removed. The dictionary always represents a sampled subset of all data, and the extra computation to maintain this dictionary is manageable. In conjunction with this, it is beneficial to remove the oldest mini-batch as it has the most outdated features and is therefore the most inconsistent with the latest key.

Weighted contrastive loss. We use the cosine distance to calculate the similarity s of features between mini-batch and memory bank. Moreover, given a query/anchor sample, we split the samples to L negative pairs and K positive pairs according to the pseudo labels, where weighted contrastive loss [36] is suitable for this issue. Thus, weighted contrastive loss is proposed to better calculate the loss in terms of multiple positive samples, which is beneficial for the optimization of model:

$$\mathcal{L}_{WCL} = \log[1 + \sum_{j=1}^{L} \exp\left(\gamma \alpha_j^n \left(s_j^n - m\right)\right)$$

$$\sum_{j=1}^{P} \exp\left(-\gamma \alpha_k^p \left(s_k^p - 1 + m\right)\right)],$$
(10)

where  $\alpha_j^n = [m+s_j^n]_+$  and  $\alpha_k^p = [1+m-s_k^p]_+$  are nonnegative weighting factors respectively,  $[\cdot]_+$  is the "cut-off at zero" operation, m refers to a margin in order to better separate similarity,  $\gamma$  is a scale factor, and L and P denote the number of negative pairs and positive pairs in the memory bank for the sample  $x_i$  of mini-batch.

We integrate the losses mentioned above. The total loss could be formulated as:

$$\mathcal{L}_{target} = \lambda_{tri} \mathcal{L}_{TRI} + \lambda_{a} \mathcal{L}_{G} + \lambda_{wcl} \mathcal{L}_{WCL}. \tag{11}$$

where  $\lambda_{tri}$ ,  $\lambda_g$ , and  $\lambda_{wcl}$  are weighting factors.

# 4. Experiments

#### 4.1. Datasets

We mainly evaluate our method on three person ReID datasets, including DukeMTMC-reID [34], Market-1501 [54] and MSMT17 [40]. DukeMTMC-reID [34] consists of 36,411 images which include 702 identities for training and 702 identities for testing, where all images are captured from 8 cameras. Market-1501 [54] contains 12,936 images of 751 identities for training and 19,281 images of 750 identities for testing, which are captured by 6 cameras. MSMT [40] contains 126,441 images of 4101 identities, of which 1041 identities are used for training, and these images are captured by 15 cameras. We adopt Mean average precision (mAP) and CMC top-1/5/10 accuracy as evaluation.

**Table 1.** Performance comparison to the unsupervised domain adaptative person Re-ID state-of-the-art methods on DukeMTMC-reID [34], Market-1501 [54] and MSMT17 [40] datasets. "BOT" denotes to "bag of tricks" method [31], which is a strong baseline in the ReID task.

Methods	DukeMTMC→Market1501				Market1501→DukeMTMC			
	mAP	top-1	top-5	top-10	mAP	top-1	top-5	top-10
UMDL [32] (CVPR'16)	12.4	34.5	52.6	59.6	7.3	18.5	31.4	37.6
PTGAN [40] (CVPR'18)	-	38.6	-	66.1	-	27.4	-	50.7
PUL [11] (TOMM'18)	20.5	45.5	60.7	66.7	16.4	30.0	43.4	48.5
TJ-AIDL [38] (CVPR'18)	26.5	58.2	74.8	81.1	23.0	44.3	59.6	65.0
SPGAN [9] (CVPR'18)	22.8	51.5	70.1	76.8	22.3	41.1	56.6	63.0
ATNet [26](CVPR'19)	25.6	55.7	73.2	79.4	24.9	45.1	59.5	64.2
SPGAN+LMP [10](CVPR'18)	26.7	57.7	75.8	82.4	26.2	46.4	62.3	68.0
CFSM [5] (AAAI'19)	28.3	61.2	-	-	27.3	49.8	-	-
BUC [25] (AAAI'19)	38.3	66.2	79.6	84.5	27.5	47.4	62.6	68.4
ECN [55] (CVPR'19)	43.0	75.1	87.6	91.6	40.4	63.3	75.8	80.4
UCDA [33] (ICCV'19)	30.9	60.4	-	-	31.0	47.7	-	-
PDA-Net [22] (ICCV'19)	47.6	75.2	86.3	90.2	45.1	63.2	77.0	82.5
PCB-PAST [51] (ICCV'19)	54.6	78.4	-	-	54.3	72.4	-	-
SSG [42] (ICCV'19)	58.3	80.0	90.0	92.4	53.4	73.0	80.6	83.2
ACT [41] (AAAI'20)	60.6	80.5	-	-	54.5	72.4	-	-
MPLP+MMCL [37] (CVPR'20)	60.4	84.4	92.8	95.0	51.4	72.4	82.9	85.0
DAAM [18] (AAAI'20)	48.8	71.3	82.4	86.3	53.1	77.8	89.9	93.7
AD-Cluster [45] (CVPR'20)	68.3	86.7	94.4	96.5	54.1	72.6	82.5	85.5
MMT [14] (ICLR'20)	71.2	87.7	94.9	96.9	65.1	78.0	88.8	92.5
NRMT [53](ECCV'20)	71.7	87.8	94.6	96.5	62.2	77.8	86.9	89.5
B-SNR+GDS-H [20](ECCV'20)	72.5	89.3	-	-	59.7	76.7	-	-
MEB-Net [46](ECCV'20)	76.0	89.9	96.0	97.5	66.1	79.6	88.3	92.2
Source Pretrain	33.9	65.3	79.2	84.0	35.2	53.3	67.7	73.1
Our GLT	79.5	92.2	96.5	97.8	69.2	82.0	90.2	92.8
Supervised learning (BOT [31])	85.7	94.1	-	-	75.8	86.2	-	-
Methods	Marke1501→MSMT17			DukeMTMC→MSMT17				
	mAP	top-1	top-5	top-10	mAP	top-1	top-5	top-10
PTGAN [40] (CVPR'18)	2.9	10.2	-	24.4	3.3	11.8	-	27.4
ENC [55] (CVPR'19)	8.5	25.3	36.3	42.1	10.2	30.2	41.5	46.8
SSG [42] (ICCV'19)	13.2	31.6	-	49.6	13.3	32.2	-	51.2
DAAM [18] (AAAI'20)	20.8	44.5	-	-	21.6	46.7	-	-
MMT [14] (ICLR'20)	22.9	49.2	63.1	68.8	23.3	50.1	63.9	69.8
Source Pretrain	7.3	19.8	29.4	34.5	11.2	31.5	42.3	47.7
Our GLT	26.5	56.6	67.5	72.0	27.7	59.5	70.1	74.2
Supervised learning (BOT [31])	48.3	72.3	-	-	48.3	72.3	-	-

#### 4.2. Implementation Details

**Training data organization.** For our joint training strategy, each mini-batch contains 64 source-domain images of 4 ground-truth identities (16 for each identity) and 64 target-domain images of 4 pseudo identities. The pseudo identities are assigned by the clustering algorithm. All images are resized to  $256 \times 128$ , and random perturbations are applied to each image, *e.g.* randomly erasing, cropping and flipping.

**Hyper-parameters.** We tune the hyper-parameters of the proposed framework on the task of Market $\rightarrow$ Duke, and the chosen hyper-parameters are directly applied to all the other tasks. ADAM optimizer is adopted to optimize the networks with weighting factors  $\lambda_{tri}=1, \lambda_g=1, \lambda_{mcl}=0.05$  and the triplet margin m=0.3. The initial learning rates (lr) are set to 0.00035. The source-domain pre-training iterates for 30 epochs and the learning rate decreases to 1/10 of its previous value every 10 epochs. We use the K-means to initialize the pseudo label, and we utilize the group-aware

label transfer to refine the pseudo label per epoch. We set  $k=\{500,1000,1500,2000\}$  for all datasets in the K-means setting.

#### 4.3. Comparison with the State-of-the-art Methods

We compare our proposed GLT with the state-of-theart methods on four domain adaptation settings in Tab. 1, *i.e.* Duke $\rightarrow$ Market, Market $\rightarrow$ Duke, Market $\rightarrow$ MSMT, and Duke $\rightarrow$ MSMT. Specifically, we surpass the clustering-based SSG [42] by considerable margins of 21.2%, 15.8%, 13.3%, and 14.4% improvements in terms of mAP on these four tasks with simpler network architectures and no extra computation of local features. DAAM [18] also adopts the clustering-based pipeline with the guidance of attention mechanism and our method shows a noticeable 30.7%, 15.1%, 5.7% and 6.1% improvements in terms of mAP on these four tasks without the help of attention mechanism. Meanwhile, our GLT adopts the one model to

significantly surpass the two-model methods MMT [14] that uses the same backbone, showing a noticeable 8.3%, 4.1%, 3.6% and 4.4% improvements in terms of mAP. Our GLT significantly outperforms the second best UDA method MEB-Net by 3.5%and3.1% in mAP accuracy, for Duke→Market, Market→Duke, respectively. Moreover, without any manual annotations, our unsupervised domain adaptive method significantly bridges the gap between unsupervised and supervised performance on person re-identification, i.e., 92.2% vs. 94.1% top-1 accuracy in Duke→Market task [31].

#### 4.4. Ablation Studies

In this section, we evaluate each components of our proposed framework by conducting ablation studies on Duke—Market and Market—Duke tasks. The experimental results are shown in Tab. 2.

Effectiveness of proposed components. We conduct extensive ablation studies to investigate the effectiveness of the proposed components in Tab. 2, i.e., label transfer, group-aware label transfer, and target instance memory bank. When introducing the label transfer algorithm, the performance is improved by 7.8% rank-1 accuracy and 6.9% mAP on Duke→Market task. By sensibly adopting memory bank with weighted contrastive loss, better representation learning enables the model to increase the rank-1 accuracy from 88.1% to 89.0% and mAP from 66.4% to 68.2%, respectively. By taking account of the group-aware label transfer, the performance is improved by 11.3% rank-1 accuracy and 3.0% mAP, which indicates that the groupaware label transfer can provides good latent pseudo-label groups for improving the quality of representation learning. In addition, group-aware strategy without the guidance of label transfer bring a little improvement due to the lack of label transfer for pseudo label refinery. This illustrates the importance of group-aware label transfer module, not just group-aware strategy. These experimental results prove the necessity and effectiveness of our group-aware label transfer for correcting noise pseudo labels.

Effectiveness of loss functions. We validate the effectiveness of multi-group ID loss, weighted contrastive loss and triplet loss in our proposed GLT framework. Triplet loss brings 3.2% and 2.1% improvements in terms of mAP, and weighted contrastive loss brings 2.6% and 2.8% improvements in terms of mAP in the Market→Duke and Duke→Market tasks, respectively. Triplet loss utilizes the hardest information within a mini-batch of instance pairs in each iteration. Meanwhile, Weighted contrastive loss in the memory bank mines the global negative and positive instances among the samples of whole dataset. The both losses can help each other.

**Table 2.** Ablation studies for our proposed framework on individual components. **TMB**: target instance memory bank. **G**: baseline with the multi-group pseudo labels. **LT**: label transfer. **GLT**: group-aware label transfer.

Methods	Duke→Market					
Wethods	mAP	top-1	top-5	top-10		
Supervised learning	85.7	94.1	-	-		
Source Pretrain	38.5	65.7	79.2	84.0		
Baseline	59.5	80.3	90.4	93.0		
Baseline + G	63.0	84.7	91.4	93.7		
Self-labeling [1]	40.9	66.6	80.6	86.3		
Baseline + TMB	62.4	82.1	91.8	94.9		
Baseline + LT	66.4	88.1	94.4	96.2		
Baseline + TMB + LT	68.2	89.0	94.5	96.1		
Baseline + G + TMB	65.2	84.3	94.8	96.7		
Baseline + G + GLT	78.7	91.4	96.7	97.4		
Baseline + $G$ + $TMB$ + $GLT$	79.5	92.2	96.5	97.8		

**Table 3.** Evaluation of the effectiveness of triplet loss and weighted contrastive loss, "w/o WCL" denotes to using the weighted contrastive loss [16].

Loss Functions	Duke-	→Market	Market→Duke		
Loss Functions	mAP	top-1	mAP	top-1	
GLT w/o TRI loss	77.3	90.9	67.1	81.2	
GLT w/o WCL loss	77.9	90.8	66.4	80.7	
GLT	79.5	92.2	69.2	82.0	

#### 5. Conclusion

To solve the problem of UDA person ReID, we propose a novel framework, the group-aware label transfer method (GLT), to combine the accurate pseudo-label prediction and effective ReID representation learning in one unified optimization objective. The holistic and immediate interaction between these two steps in the training process can greatly help the UDA person ReID task. To achieve this goal, we consider the online label refinement problem as an optimal transport problem, which explores the minimal cost of assigning M samples to N pseudo labels. More importantly, we introduce a group-aware strategy to assign implicit attribute group IDs to samples. The combination of the online label refining algorithm and the group-aware strategy can better correct the noisy pseudo label in an online fashion and narrow down the search space of the target identity. Our method not only achieves state-of-the-art performance but also significantly closes the gap between supervised and unsupervised performance on person re-identification.

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