



Graph Sampling Based Deep Metric Learning for Generalizable Person Re-Identification

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Abstract

Recent studies show that, both explicit deep feature matching as well as large-scale and diverse training data can significantly improve the generalization of person reidentification. However, the efficiency of learning deep matchers on large-scale data has not yet been adequately studied. Though learning with classification parameters or class memory is a popular way, it incurs large memory and computational costs. In contrast, pairwise deep metric learning within mini batches would be a better choice. However, the most popular random sampling method, the well-known PK sampler, is not informative and efficient for deep metric learning. Though online hard example mining has improved the learning efficiency to some extent, the mining in mini batches after random sampling is still limited. This inspires us to explore the use of hard example mining earlier, in the data sampling stage. To do so, in this paper, we propose an efficient mini-batch sampling method, called graph sampling (GS), for large-scale deep metric learning. The basic idea is to build a nearest neighbor relationship graph for all classes at the beginning of each epoch. Then, each mini batch is composed of a randomly selected class and its nearest neighboring classes so as to provide informative and challenging examples for learning. Together with an adapted competitive baseline, we improve the state of the art in generalizable person re-identification significantly, by 25.1% in Rank-1 on MSMT17 when trained on RandPerson. Besides, the proposed method also outperforms the competitive baseline, by 6.8% in Rank-1 on CUHK03-NP when trained on MSMT17. Meanwhile, the training time is significantly reduced, from 25.4 hours to 2 hours when trained on RandPerson with 8,000 identities. Code is available at https://github.com/ShengcaiLiao/QAConv.

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1. Introduction

Person re-identification is a popular computer vision task, where the goal is to find a person, given in a query image, from the search over a large set of gallery images. In the last two years, generalizable person re-identification has gain increasing attention due to both its research and practical value [11, 12, 17, 21, 26, 46, 48]. This task studies the generalizability of a learned person re-identification model in unseen scenarios, and employs direct cross-dataset evaluation [10, 38] for performance benchmarking.

For deep metric learning, beyond feature representation learning and loss designs, explicit deep feature matching schemes are shown to be effective for matching person images [1, 14, 17, 24, 28], due to the advantages in addressing pose and viewpoint changes, occlusions, and misalignments. In particular, a recent method, called query-adaptive convolution (QAConv) [17], has proved that explicit convolutional matching between gallery and query feature maps is quite effective for generalizable person re-identification. However, these methods all require more computational costs compared to conventional feature learning methods.

Beyond novel generalizable algorithms, another way to improve generalization is to enlarge the scale and diversity of the training data. For example, a recent dataset called RandPerson [33] synthesized 8,000 identities, while [31] and [2] both collected 30K persons for re-identification training. These studies all observed improved generalization ability for person re-identification. However, the efficiency of deep metric learning from large-scale data has not yet been adequately studied in person re-identification.

There are some popular ways of learning deep person re-identification models, including classification (with the ID loss [43]), metric learning (with a pairwise loss [5, 38] or triplet loss [9]), and their combinations (e.g. ID + triplet loss). Using an ID loss is convenient for classification learning. However, in large-scale deep learning, involving classifier parameters incurs large memory and computational costs in both the forward and backward passes. Similarly,

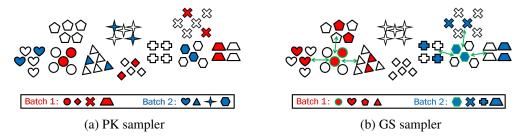


Figure 1. Two different sampling methods: (a) PK sampler; and (b) the proposed GS sampler. Different shapes indicate different classes, while different colors indicate different batches. GS constructs a graph for all classes and always samples nearest neighboring classes.

involving class signatures for metric learning in a global view is also not efficient. For example, QAConv in [17] is difficult to scale up for large-scale training, because a class memory module is designed, where full feature maps are stored for all classes as signatures, and they are required for cross feature map convolutional matching during training.

Therefore, involving class parameters or signatures in either classification or metric learning is not efficient for large-scale person re-identification training. In contrast, we consider that pairwise deep metric learning between samples in mini batches is better suited for this task. Accordingly, the batch sampler plays an important role for efficient learning [9, 37]. The well-known PK sampler [9, 22] is the most popular random sampling method in person reidentification. It first randomly selects P classes, and then randomly samples K images per class to construct a mini batch of size $B = P \times K$. Since this is performed randomly, the sampled instances within a mini batch are uniformly distributed across the whole dataset (see Fig. 1 (a)), and might therefore not be informative and efficient for deep metric learning. To address this, an online hard example mining method was proposed in [9], which improved the learning efficiency to some extent. However, the mining is performed online on the already sampled mini batches. Therefore, this method is still limited by the fully random PK sampler, because the mini batches obtained by this sampler do not consider the sample relationship information.

To address this, we propose to shift the hard example mining earlier to the data sampling stage. Accordingly, we propose an efficient mini-batch sampling method, called graph sampling (GS), for large-scale deep metric learning. The basic idea is to build a nearest neighbor relationship graph for all classes at the beginning of each epoch. Then, the mini-batch sampling is performed by randomly selecting a class as anchor, and its top-k nearest neighboring classes, with the same K instances per class, as shown in Fig. 1 (b). This way, instances within a sampled mini batch are mostly similar to each other, so as to provide informative and challenging examples for discriminant learning. From face recognition loss function studies [4,19,35], it is known that focusing on boundary (hard) examples helps improving

the discriminant ability of the learned model, and helps resulting in compact representations that generalize well beyond the training data. The GS sampler shares a similar idea in focusing on nearest neighboring classes, and thus has a potential of improving the discrimination and generalization ability of the learned model.

In summary, the contributions of this paper include: (1) We propose a new mini-batch sampling method, termed GS, and prove that it enables more efficient learning than the well-known PK sampler; (2) We improve a very competitive baseline by 6.8% in Rank-1 with MSMT17 → CUHK03-NP, and reduce the training time significantly, from 25.4 hours to 2 hours on RandPerson with 8,000 identities; and (3) Together with the baseline, we improve the state of the art in generalizable person re-identification significantly, by 20.6% in Rank-1 with Market-1501 → MSMT17, and by 25.1% in Rank-1 with RandPerson → MSMT17.

2. Related Work

Metric learning approaches have been widely studied in the early stage of person re-identification. Many algorithms have been proposed, such as the well-known PRDC [44], KISSME [13], and XQDA [16], to name a few. In recent years, deep metric learning in particular has become popular and been extensively studied. Beyond feature representation learning, specific deep metric learning can be roughly classified in terms of loss function designs and deep feature matching schemes. For loss function designs, pairwise loss functions [5, 38], classification or identification loss [43], and triplet loss [9, 22, 44] are the most popular. For deep feature matching schemes, a number of methods have been proposed in the literature. For example, Ahmed et al. proposed a deep convolutional architecture with layers specifically designed for local neighborhood matching [1]. Li et al. proposed a novel filter pairing neural network (FPNN) to jointly handle several known challenges, such as misalignment and occlusions [14]. Shen et al. proposed an end-to-end deep Kronecker-Product Matching (KPM) network [24] for softly aligned matching. Suh et al. proposed a deep neural network to learn part-aligned bilinear representations [28]. Liao and Shao proposed the query-adaptive

convolution (QAConv) for explicit deep feature matching, and proved its effectiveness for generalizable person reidentification [17]. They also proposed a Transformer based method, TransMatcher [18], for improved performance.

Generalizable person re-identification was initially studied in [10, 38], where direct cross-dataset evaluation was proposed to benchmark algorithms. With advancements in deep learning, this task has gained increasing attention in recent years. For example, Song et al. [26] proposed a domain-invariant mapping network with a meta-learning pipeline. Jia et al. [11] adopted both instance and feature normalization to alleviate both style and content variances across datasets. Zhou et al. proposed a new backbone network called OSNet [46], and further demonstrated its advantages in generalizing deep models [46]. Jin et al. proposed a style normalization and restitution module, which shows good generalizability [12]. Yuan et al. proposed an adversarial domain-invariant feature learning network (ADIN), which explicitly learns to separate identityrelated features from challenging variations [39]. Zhuang et al. proposed a camera-based batch normalization (CBN) method for domain-invariant representation learning [48]. Recently, meta-learning has also been shown to be effective for learning generalizable models. For example, Zhao et al. proposed memory-based multi-source meta-learning (M³L) for generalizing to unseen domains [41]. Choi et al. proposed the MetaBIN algorithm for meta-training the batch-instance normalization nettwork [3]. Bai et al. proposed a dual-meta generalization network and a large-scale dataset called Person30K for person re-identification [2]. In addition to the above, Wang et al. proposed a large-scale synthetic person re-identification dataset, called RandPerson, and proved that models learned from synthesized data generalize well to real-world datasets [33].

However, the generalization of current methods is still far from satisfactory for practical person re-identification. Taking face recognition as a good example in practice, future directions may gradually be learning from more large-scale data for better performance. However, the efficiency of large-scale learning has been inadequately studied in person re-identification. As basic as the mini-batch sampler, though it plays an important role in deep metric learning [9, 36, 37], it still has not yet been much studied.

Beyond online hard example mining within mini batches [9], several methods have been proposed for hard example mining during data sampling for mini batches. Suh et al. [27] proposed a stochastic class-based hard example mining for deep metric learning. It uses learnable class signatures to find nearest classes, and further performs an instance-level refined search within the subset of classes found in the first stage for hard example mining. Besides, the Doppelganger [25] also relies on classification layers for doppelganger mining from the predicted classification

scores. However, these methods require classification parameters to be learned for class mining, which is intractable for large-scale classes and complex non-Euclidean matchers (e.g. QAConv). In [30], all training classes are divided into subspaces by clustering on averaged class representations, and then mini batches are sampled within each subspace. This method requires a full forward pass of all the training data, and the clustering operation cannot easily be scaled up to large-scale classes. In [7], SmartMining was proposed, which builds an approximate nearest neighbor graph for all training samples after a full forward pass of the training data for feature extraction. However, this instance-level mining can be very expensive in computation, and even infeasible for complex non-Euclidean metric layers. In contrast, we propose and prove that sampling one example per class for class mining works well for large-scale deep metric learning without classification or instance-level mining.

3. Deep Metric Learning

There are two popular ways for learning deep person reidentification neural networks. The first one is the classification based method [43], also known as using the identification loss, or ID loss. This is a straightforward extension from general image classification. Since person reidentification is an open-class problem, the learned classifier is usually dropped after training. The last feature embedding layer is usually adopted instead (known as the identity embedding, or IDE [43]), and the Euclidean or cosine distance is applied to measure the distance between two person images. The second one is the triplet loss based method [9,23], which is usually combined with the ID loss. Together with the online hard example mining, the triplet loss is a very useful auxiliary loss function for enhancing the discriminability of the learned model.

However, the above methods always require classifier parameters, which incur large memory and computational costs in both the forward and backward passes of large-scale deep learning. When dot products are employed for classification this is still acceptable to some extent. However, with more complex modules, e.g. QAConv [17] where a full feature map convolution is required for matching, learning with class signatures is difficult to scale up.

Therefore, for large-scale deep metric learning, we consider removing classification layers. Accordingly, pairwise matching between mini-batch samples is another solution [15,38]. We adopt QAConv as our baseline method, which is the recent state of the art for generalizable person reidentification. It constructs query adaptive convolutional kernels on the fly for image matching, which suits pairwise learning. However, the original design of QAConv learning is based on the so-called class memory, which stores one feature map for each class for image-to-class matching, instead of using pairwise matching between mini-batch sam-

ples. Considering the matching complexity of the QAConv layer, this is not efficient in large-scale learning. Therefore, we only consider pairwise matching between mini-batch samples for QAConv, and remove its class memory.

4. Graph Sampling

4.1. Motivation

As discussed, for deep metric learning, the well-known PK sampler [9] is typically used to provide mini-batch samples. However, its random nature makes the sampled instances not informative enough for discriminant learning. In the PK sampler, as shown in Fig. 1 (a), P classes and K images per class are randomly sampled for each mini batch. Though an online hard example mining (OHEM) was further proposed in [9] to find informative instances within a mini batch, the PK sampler itself is still not efficient, as it provides limited hard examples for OHEM to mine.

Therefore, the sampling method itself needs to be improved so as to provide informative samples for mini batches. Instead of using fully random sampling, the relationships among classes need to be considered. Thus, we construct a graph for all classes at the beginning of each epoch, and always sample nearest neighboring classes in a mini batch so as to enable discriminant learning. We call this idea graph sampling (GS), which is detailed below.

4.2. GS Sampler

At the beginning of each epoch, we utilize the latest learned model to evaluate the distances or similarities between classes, and then construct a graph for all classes. This way, the relationships between classes can be used for informative sampling. Specifically, we randomly select one image per class to construct a small sub-dataset. Then, the feature embeddings with the current network are extracted, denoted as $\mathbf{X} \in R^{C \times d}$, where C is the total number of classes for training, and d is the feature dimension. Next, pairwise distances between all the selected samples are computed, e.g. by QAConv. As a result, a distance matrix $dist \in R^{C \times C}$ for all classes is obtained.

Then, for each class c, the top P-1 nearest neighboring classes can be retrieved, denoted by $\mathcal{N}(c)=\{x_i|i=1,2,\ldots,P-1\}$, where P is the number of classes to sample in each mini batch. Accordingly, a graph G=(V,E) can be constructed, where $V=\{c|c=1,2,\ldots,C\}$ represents the vertices, with each class being one node, and $E=\{(c_1,c_2)|c_2\in\mathcal{N}(c_1)\}$ represents the edges.

Finally, for the mini-batch sampling, for each class c as anchor, we retrieve all its connected classes in G. Then, together with the anchor class c, we obtain a set $A = \{c\} \bigcup \{x | (c,x) \in E\}$, where |A| = P. Next, for each class in A, we randomly sample K instances per class to generate a mini batch of $B = P \times K$ samples for training. A

pseudocode of the GS sampler is shown in Appendix A.

Note that, different from other mini-batch sampling methods, for the GS sampler the number of mini batches or iterations per epoch is always C, which is independent to the parameters B, P, and K. Nevertheless, the parameter B still affects the computational load of each mini batch. Besides, one may worry that the GS sampler will be computationally expensive. However, note that, firstly, only one image per class is randomly sampled for the graph construction; and, secondly, the above computation is performed only once per epoch. In practice, we find that the GS sampler with QAConv, which is already a heavy matcher compared to the mainstream Euclidean distance, only requires tens of seconds for thousands of identities. Details will be presented in the experimental section.

4.3. Loss Function

With mini batches provided by the GS sampler, we apply QAConv to compute similarity values between each pair of images, and formulate a triplet-based ranking learning problem within mini batches. Accordingly, we compute the batch OHEM triplet loss [9] alone for metric learning:

$$\ell(\boldsymbol{\theta}; X) = \sum_{i=1}^{P} \sum_{a=1}^{K} [m - \min_{\substack{p=1...K}} s(f_{\boldsymbol{\theta}}(x_i^a), f_{\boldsymbol{\theta}}(x_i^p)) + \max_{\substack{j=1...P\\j \neq i\\n=1...K}} s(f_{\boldsymbol{\theta}}(x_i^a), f_{\boldsymbol{\theta}}(x_j^n))]_+,$$
(1)

where $X = \{x_i^a, i \in [1, P], a \in [1, K]\}$ contains the minibatch samples, $\boldsymbol{\theta}$ is the network parameter, $f_{\boldsymbol{\theta}}$ is the feature extractor, $s(\cdot, \cdot)$ is the similarity, and m is the margin.

Note that Eq. (1) is usually used as an auxiliary to the ID loss, but not alone in person re-identification. This is probably because random samplers including PK cannot provide informative mini batches for OHEM to mine, which makes Eq. (1) very small or even zero, and so the learning is not efficient. In contrast, with the proposed GS sampler, we prove that the OHEM triplet loss works well by itself.

4.4. Gradient Clipping

Note that the GS sampler already provides almost the hardest mini batches, and the batch OHEM triplet loss further finds the hardest triplets within a mini batch for training. As a result, the model may suffer optimization difficulty, which in turn may impact convergence during training. In practice, we find that limiting K=2 alleviates this problem significantly. Or otherwise, the binary crossentropy loss for pairwise matching can be a more stable alternative to the OHEM triplet loss (see Appendix B).

Furthermore, to stabilize the training with the GS sampler and the hard triplet loss, we clip the gradient norm during the backward propagation. Specifically, let g be the gradient of all parameters, and $\|\mathbf{g}\|$ be its norm. The gradient

will be clipped as $\mathbf{g} \leftarrow \min(1, \frac{T}{\|\mathbf{g}\|}) \cdot \mathbf{g}$, where T is a predefined threshold. That is, if the gradient norm is larger than T then clip it to be T. Note that GS and OHEM provide the hardest examples, which facilitates discriminant learning. However, this may also lead to overfitting. Therefore, besides stabilizing the training, the gradient clipping operation is also useful to regularize noisy gradients to avoid overfitting on source domain, and, in turn, improving the generalization performance. The effect of this gradient clipping will be analyzed in the experiments.

5. Experiments

5.1. Implementation Details

Our implementation is adapted from the official PyTorch code of QAConv [17] (MIT license). We first build an improved baseline based on QAConv. Specifically, ResNet-50 [8] is used as the backbone, with IBN-b layers appended, following several recent studies [11, 12, 20, 46, 48]. The layer3 feature map is used, with a neck convolution of 128 channels appended as the final feature map. The input image size is 384×128 (see Appendix E for results with 256×128). Several commonly used data augmentation methods are applied, including random cropping, flipping, occlusion [17], and color jittering. The batch size is set to 64. The SGD optimizer is adopted to train the network, with a learning rate of 0.0005 for the backbone, and 0.005 for newly added layers. The maximal learning epochs are 60. When the initial loss is reduced as a factor of 0.7, the learning rates are decayed by 0.1, and an early stopping is triggered after a further half of the already learned epochs. Gradient clipping is applied with T=8. Automatic Mixed Precision (AMP) in PyTorch is applied to accelerate training. When the proposed GS sampler is further applied (denoted by QAConv-GS), we use the hard triplet loss (m=16), instead of the class memory based loss proposed in [17], and the default parameters for GS are B=64, and K=2.

5.2. Datasets

Four large-scale person re-identification datasets, CUHK03 [14], Market-1501 [42], MSMT17 [34], and RandPerson [33] are used in our experiments. The CUHK03 dataset contains 1,360 persons and 13,164 images. The most challenging subset named detected is used for our experiments. Besides, the CUHK03-NP protocol [45] is adopted, with 767 and 700 subjects used for training and testing, respectively. The Market-1501 dataset includes 32,668 images of 1,501 identities captured from six cameras. The training subset contains 12,936 images from 751 identities, while the test subset includes 19,732 images from 750 identities. The MSMT17 dataset contains 4,101 identities and 126,441 images captured from 15 cameras. It is divided into a training set of 32,621 images

from 1,041 identities, and a test set with the remaining images from 3,010 identities. The RandPerson dataset is a recently released synthetic person re-identification dataset. It contains 8,000 persons and 1,801,816 images. We use the subset including 132,145 images of the 8,000 identities. This dataset is only used for large-scale training and generalization testing. Cross-dataset evaluation [10,38] is performed on all datasets, by training on the training subset of one dataset (except that with MSMT17 we further used an additional setting with all images for training), and evaluating on the test subset of another dataset. Rank-1 and mean average precision (mAP) are used as the performance metrics, evaluated under single-query evaluation protocol.

5.3. Comparison to the State of the Art

A comparison to the state of the art (SOTA) in generalizable person re-identification is shown in Table 1, where three datasets are used for training, and three others are used for testing. Note that, with MSMT17 as the training set, one setting is to use all images for training, regardless of its subset splits. This is denoted by MSMT17 (all). Several generalizable person re-identification methods published recently are compared, including OSNet-IBN [46], OSNet-AIN [47], MuDeep [21], SNR [12], QAConv [17], CBN [48], ADIN [39], and M³L [41]. Table 1 shows that QAConv-GS significantly improves the previous SOTA. For example, with Market-1501 → CUHK03, the Rank-1 and mAP are improved by 8.8% and 9.0%, respectively. With Market-1501 \rightarrow MSMT17, they are improved by 20.6% and 7.7%, respectively. With MSMT17 (all) \rightarrow Market-1501, the improvements are 9.8% for Rank-1 and 13.8% for mAP. With RandPerson as the training data, the improvements on Market-1501 are 12% for Rank-1 and 7.4% for mAP, while the improvements on MSMT17 are 25.1% for Rank-1 and 8.7% for mAP. Though RandPerson is synthetic, the results show that models learned on it generalize quite well to realworld datasets, which confirms the findings in [33].

Note that, M³L [41] uses a different evaluation protocol, and thus the results are not directly comparable. Specifically, M³L is trained on three datasets selected from CUHK03, Market-1501, DukeMTMC-reID¹, and MSMT17, while the other is held for testing. Impressive results are obtained by M³L on CUHK03-NP, which, though not directly comparable, exceed all our results, including those trained with all MSMT17 images. However, on Market-1501, the proposed method trained on MSMT17 outperforms M³L in Rank-1 by 3.2%, while the mAPs are comparable. Furthermore, on MSMT17, the proposed method trained on Market-1501 significantly outperforms M³L, with 9% gain in Rank-1 and 2.5% in mAP. This is quite encouraging, since in both cases our training dataset is a subset of that used by M³L.

¹It is no longer available, so we do not use it in our experiments.

Method	Venue	Training	CUHK)3-NP	Market-1501		MSMT17	
		Training	Rank-1	mAP	Rank-1	mAP	Rank-1	mAP
M^3L [41]	CVPR'21	Multi	33.1	32.1	75.9	50.2	36.9	14.7
MGN [21, 32]	ACMMM'18	Market-1501	8.5	7.4	95.7	86.9	-	-
MuDeep [21]	TPAMI'20	Market-1501	10.3	9.1	95.3	84.7	-	-
QAConv [17]	ECCV'20	Market-1501	9.9	8.6	-	-	22.6	7.0
OSNet-AIN [47]	TPAMI'21	Market-1501	-	-	94.2	84.4	23.5	8.2
CBN [48]	ECCV'20	Market-1501	-	-	91.3	77.3	25.3	9.5
QAConv-GS	Ours	Market-1501	19.1	18.1	91.6	75.5	45.9	17.2
PCB [29, 39]	ECCV'18	MSMT17	-	-	52.7	26.7	-	-
MGN [32, 39]	ACMMM'18	MSMT17	-	-	48.7	25.1	-	-
ADIN [39]	WACV'20	MSMT17	-	-	59.1	30.3	-	-
SNR [12]	CVPR'20	MSMT17	-	-	70.1	41.4	-	-
CBN [48]	ECCV'20	MSMT17	-	-	73.7	45.0	72.8	42.9
QAConv-GS	Ours	MSMT17	20.9	20.6	79.1	49.5	79.2	50.9
OSNet-IBN [46]	CVPR'19	MSMT17 (all)	-	-	66.5	37.2	-	-
OSNet-AIN [47]	TPAMI'21	MSMT17 (all)	-	-	70.1	43.3	-	-
QAConv [17]	ECCV'20	MSMT17 (all)	25.3	22.6	72.6	43.1	-	-
QAConv-GS	Ours	MSMT17 (all)	27.6	28.0	82.4	56.9	-	-
RP Baseline [33]	ACMMM'20	RandPerson	13.4	10.8	55.6	28.8	20.1	6.3
CBN [40]	ECCV'20	RandPerson	-	-	64.7	39.3	20.0	6.8
QAConv-GS	Ours	RandPerson	18.4	16.1	76.7	46.7	45.1	15.5

Table 1. Comparison of the state-of-the-art direct cross-dataset evaluation results (%). MSMT17 (all) means all images are used for training, regardless of subset splits. M³L is trained on three datasets selected from CUHK03, Market-1501, DukeMTMC-reID, and MSMT17, while the other is held for testing. Results in gray cells are with within-dataset evaluation for a reference. "-" means not reported or not applicable.

5.4. Ablation Study

5.4.1 Comparison of QAConv variants

Table 2 shows a comparison among different variations of QAConv: the original QAConv [17] (denoted as Ori), the competitive QAConv baseline we adapted (denoted as Base), and the proposed QAConv-GS. It shows that, beyond the successful learning scheme of the class memory module proposed in QAConv, QAConv-GS with the proposed GS sampler is also very effective in learning discriminant models. QAConv-GS outperforms the competitive baseline for all experiments, by 6.8% and 5.4% in Rank-1, respectively, on CUHK03 and Market-1501 when trained on MSMT17.

Interestingly, Table 2 also shows that the within-dataset evaluation results are also improved by QAConv-GS compared to the baseline. However, improving performance on a single dataset does not always lead to better generalization, since it may also overfit a dataset, as can be observed in Table 1. Therefore, we suggest a focus on generalization since it is more critical for practical applications.

Furthermore, we also compare the training time of QA-Conv (with class memory) and QAConv-GS. Both methods are tested on a single NVIDIA V100 GPU. From the comparison shown in Table 2, it can be observed that the original QAConv learned with class memory becomes very

	Training		CUHK03		Market		MSMT17	
	Data	Hours	R1	mAP	R1	mAP	R1	mAP
Ori	Market	1.33	9.9	8.6	-	-	22.6	7.0
Base	Market	0.47	14.6	14.6	88.7	71.4	42.6	15.8
GS	Market	0.25	19.1	18.1	91.6	75.5	45.9	17.2
Base	MSMT	1.33	14.1	15.7	73.7	44.7	72.5	43.4
GS	MSMT	0.73	20.9	20.6	79.1	49.5	79.2	50.9
Ori	MS-all	26.9	25.3	22.6	72.6	43.1	-	-
Base	MS-all	15.0	23.4	23.1	80.1	53.2	-	-
GS	MS-all	3.42	27.6	28.0	82.4	56.9	-	-
Base	RP	25.4	15.2	14.6	75.9	46.0	44.4	15.5
GS	RP	2.0	18.4	16.1	76.7	46.7	45.1	15.5

Table 2. Comparison of QAConv variants. Ori: the original QAConv [17]. Base: the competitive baseline we adapted. GS: graph sampling (ours). MS-all: MSMT17 (all). RP: RandPerson.

slow when trained on large-scale datasets, such as the full MSMT17 or RandPerson. This is not surprising, because in each mini-batch iteration, the QAConv with class memory needs to compute matching scores between mini-batch samples and the feature map memory of all classes; and the number of classes is 4,101 in MSMT17, and 8,000 in Rand-

Person. In contrast, the proposed pairwise learning with the GS sampler is much more efficient because it avoids matching all classes in each iteration. As can be seen from Table 2, the training time of the baseline QAConv can be reduced from 25.4 hours to 2 hours when trained on RandPerson with 8,000 identities, which is a significant achievement.

In addition, we also evaluate the sampling efficiency of the proposed GS sampler. As stated earlier, it constructs a graph at the beginning of each epoch. We evaluate the running time of all the computations in GS. The results are 4 seconds on Market-1501, 9 seconds on the MSMT17 training subset, 40 seconds on the full MSMT17 dataset, and 138 seconds on RandPerson with 8,000 identities. Therefore, the GS sampler is in fact efficient, despite being incorporated into QAConv, which is a heavy matcher compared to the mainstream Euclidean distance.

5.4.2 Comparison of different sampling methods

In Table 3, using the same QAConv and hard triplet loss, we compare three mini-batch sampling methods, including PK, a clustering based method [30] (denoted as Cluster), and GS. Besides, the implementation of PK and Cluster follows GS, where the number of batches per epoch is determined by the number of classes. For [30], since k-means does not support non-Euclidean metric, we replace it with spectral clustering. The subspace parameter M in [30] is set to 10, after an optimization in [5, 50]. From Table 3, we can see that PK performs the worst, due to its fully random nature, which does not provide enough hard examples in mini batches. Besides, we can see that, with the subspace clustering method proposed in [30], the performance is generally improved, thanks to the more informative mini batches sampled within each cluster. However, feature extraction from the whole training set and clustering of all classes are time consuming. In contrast, the proposed GS sampler is more efficient, since it only considers one example per class for the graph construction. Furthermore, GS also achieves the best performance, with improvements over Cluster of up to 4.7% in Rank-1, and 3.4% in mAP. We believe that clustering is less effective than graph based GS due to two reasons. First, only cluster centers may be surrounded by their dense neighbors, while others, especially boundary points (classes), may not be always with their full set of neighbors in the same cluster. Second, mini-batch classes need to be randomly sampled within a cluster, of which the operation may further miss out some nearest neighbors of each class.

Furthermore, in Appendix C, applications to two other baselines, OSNet [46] and TransMatcher [18], also verify the generality of GS's advantage over PK. Besides, application to unsupervised domain adaptation (UDA) with GS in SpCL [6] is discussed in Appendix D, and a variant of GS using class centers is analyzed in Appendix E.

Method	Training		CUHK03		Market		MSMT17	
Method	Data	Time	R1	mAP	R1	mAP	R1	mAP
PK	Market	99	17.9	17.0	-	-	43.3	15.6
Cluster	Market	117	18.4	17.3	-	-	44.0	15.8
GS	Market	100	19.1	18.1	-	ı	45.9	17.2
PK	MSMT	141	18.6	18.8	75.7	46.1	-	-
Cluster	MSMT	196	18.4	19.2	77.2	47.6	-	-
GS	MSMT	145	20.9	20.6	79.1	49.5	-	-
PK	MS-all	669	24.5	24.6	78.7	52.1	-	-
Cluster	MS-all	881	26.3	26.3	80.4	54.2	-	-
GS	MS-all	685	27.6	28.0	82.4	56.9	-	-
PK	RP	1,150	16.9	14.7	73.2	43.5	40.3	13.1
Cluster	RP	1,922	17.3	15.0	73.3	43.3	40.4	13.4
GS	RP	1,397	18.4	16.1	76.7	46.7	45.1	15.5

Table 3. Comparison of different sampling methods. MS-all: MSMT17 (all). RP: RandPerson. Time is with seconds per epoch.

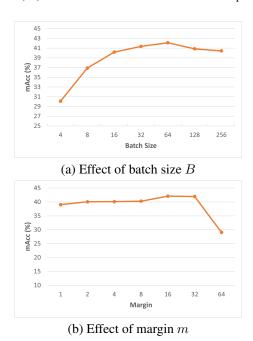


Figure 2. mAcc (%) performance with (a) different batch sizes, and (b) different margin parameters, trained on MSMT17.

5.4.3 Parameter analysis

In Fig. 2, we show the performance of the proposed method with different batch sizes and margin parameters. The training is performed on MSMT17. For ease and reliable comparison, we report the average (denoted by mAcc) of all Rank-1 and mAP results on all test sets over four random runs. We observe that, generally, the accuracy increases with increasing batch size B, but saturates at 64. As for the margin parameter m, note that the QAConv similarity score $s(\cdot, \cdot)$ used in Eq. (1) ranges in $(-\infty, +\infty)$. Fig. 2(b) shows

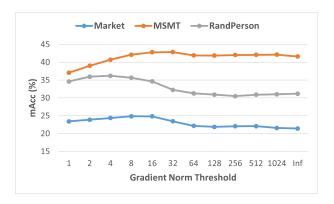


Figure 3. Influence of gradient clipping, trained on three datasets.

that the performance slightly improves with increasing m due to the increased discriminability, and achieves the best with m=16. However, after m=32, the performance drops significantly, due to intractable learning difficulty.

5.4.4 Effect of gradient clipping

Next, we study the effect of gradient clipping on the learning of QAConv-GS. The results are shown in Fig. 3. Interestingly, when trained on MSMT17, the performance is less affected without gradient clipping (Inf). Specifically, with gradient clipping, only a slight improvement can be obtained, but too small threshold T even prevent effective model learning. This is because, in our experiments, MSMT17 is the most comprehensive dataset. It provides large-scale and diverse training examples, which prevents overfitting in the view of "regularization from data". However, with the small-scale training dataset Market-1501, and the quite different synthetic dataset RandPerson, gradient clipping does provide useful regularization for model training, and improves the generalization performance. Therefore, a reasonable value of T=8 is considered as a trade-off.

5.4.5 Visualization of GS

Fig. 4 shows some examples (more in Appendix F) of the nearest neighboring classes generated by the GS sampler. As can be observed, the GS sampler is indeed able to find similar classes as hard examples to challenge the learning. For example, it identifies similar kinds of clothes, colors, patterns, and accessories. These confusing examples help a lot in learning discriminative models.

6. Conclusion

With this study, we show that the popular PK sampler is not efficient in deep metric learning, and thus we propose a new batch sampler, called the graph sampler, to help learning discriminant models more efficiently. This is achieved



Figure 4. Groups of examples of the nearest neighboring classes generated by the GS sampler when trained on (a) Market-1501 and (b) MSMT17. In each group, the upper left image is the center class, and other images are the top-7 nearest neighboring classes.

by constructing a nearest neighbor graph of all classes for informative sampling. Together with a competitive baseline, we achieve the new state of the art in generalizable person re-identification with a significant improvement. Meanwhile, the training time is much reduced by removing the classification parameters and only using the pairwise distances between mini batches for loss computation. We believe the proposed technique is general and may also be applied in other fields, such as image retrieval, and face recognition, among others. Moreover, we discuss social impacts and some limitations of this research in Appendix G and H.

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