Robust Graph Structure Learning over Images via Multiple Statistical Tests

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Abstract

Graph structure learning aims to learn connectivity in a graph from data. It is particularly important for many computer vision related tasks since no explicit graph structure is available for images for most cases. A natural way to construct a graph among images is to treat each image as a node and assign pairwise image similarities as weights to corresponding edges. It is well known that pairwise similarities between images are sensitive to the noise in feature representations, leading to unreliable graph structures. We address this problem from the viewpoint of statistical tests. By viewing the feature vector of each node as an independent sample, the decision of whether creating an edge between two nodes based on their similarity in feature representation can be thought as a *single* statistical test. To improve the robustness in the decision of creating an edge, multiple samples are drawn and integrated by *multiple* statistical tests to generate a more reliable similarity measure, consequentially more reliable graph structure. The corresponding elegant matrix form named $\mathcal{B}$**ttention** is designed for efficiency. The effectiveness of multiple tests for graph structure learning is verified both theoretically and empirically on multiple clustering and ReID benchmark datasets.

1 Introduction

Graph structure learning plays an important role in Graph Convolutional Networks (GCNs). It seeks to discover underlying graph structures for better graph representation learning when graph structures are unavailable, noisy or corrupted. These issues are typical in the field of vision tasks [54, 59, 58, 19, 40, 47, 53, 12, 37, 62, 45]. As an example, there is no explicit graph structure between photos in your phone albums, so the graph-based approaches [19, 40, 53] in photo management softwares seek to learn graph structure. A popular research direction on it is to consider graph structure learning as similarity measure learning upon the node embedding space [56]. They learn weighted adjacency matrices [19, 40, 8, 9, 41, 60, 26, 7, 24, 33] as graphs and feed them to GCNs to learn representations.

However, the main problem of the above weighted-based methods is that weights are not always accurate and reliable: two nodes with smaller associations may have a higher weight score than those with larger associations. The root of this problem is closely related to the **noise** in the high dimensional feature vectors extracted by the deep learning model, as similarities between nodes are usually decided based on their feature vectors. The direct consequence of using an inaccurate graph structure is so called feature pollution [53], leading to a performance degradation in downstream tasks. In our empirical studies, we found that close to $1/3$ of total weights are assigned to noisy edges when unnormalized cosine similarities between feature vectors are used to calculate weights.

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Many methods have been developed to improve similarity measure between nodes in graph, including adding learnable parameters [8, 26, 7], introducing nonlinearity [9, 41, 60], Gaussian kernel functions [24, 33] and self-attention [50, 11, 19, 40]. Despite the progress, they are limited to modifying similarity function, without explicitly addressing the noise in feature vectors, a more fundamental issue for graph structure learning. One way to capture the noise in feature vectors is to view the feature vector of each node as an independent sample from some unknown distribution associated with each node. As a result, the similarity score between two nodes can be thought as the expectation of a test to decide if two nodes share the same distribution. Since, only one vector is sampled for each node in the above methods, the statistical test is essentially based on a single sample from both nodes, making the calculated weights inaccurate and unreliable.

An possible solution towards addressing the noise in feature vectors is to run multiple tests, a common approach to reduce variance in sampling and decision making [4, 10, 25]. Multiple samples are drawn for each node, and each sample can be used to run an independent test to decide if two nodes share the same distribution. Based on the results from **multiple tests**, we develop a novel and robust similarity measure that significantly improves the robustness and reliability of calculated weights. This paper will show *how* to accomplish multiple tests in the absence of graph structures and theoretically prove *why* the proposed approach can produce a better similarity measure under specified conditions. This similarity measure is also crafted into an efficient and elegant form of matrix, named $\mathcal{B}$**ttention**. $\mathcal{B}$**ttention** utilizes the fourth order statistics of the features, which is a significant difference with popular attentions. Benefiting from a better similarity measure produced by multiple tests, graph structures can be significantly improved, resulting in better graph representations and therefore boosting the performance of downstream tasks. We empirically verify the effectivness of the proposed method over the task of clustering and ReID.

In summary, this paper makes the following three major contributions: (1) To the best of our knowledge, this is the first paper to address inaccurate edge weights problem by statistical tests on the task of graph structure learning over images. (2) A novel and robust similarity measure is proposed based on multiple tests, proven theoretically to be superior and designed in an elegant matrix form to improve graph structure quality. (3) Extensive experiments further verify the robustness of the graph structure learning method against noise. State-of-the-art (SOTA) performances are achieved on multiple clustering and ReID benchmarks with the learned graph structures.

2 Methodology

To handle the noise in feature vectors, we model the feature vector of each node as a sample drawn from the underlying distribution associated with that node, and the similarity score between two nodes as the expectation of a statistical test. The **test** is to compare their similarity to a given threshold to decide if two nodes share the same distribution and that decision can be captured by a binary Bernoulli random variable. The comparison in statistical test may lead to error edges, i.e., **noisy edges** (an edge connects two nodes of different categories) or **missing edges** (two nodes of the same category should be connected but are not). The superiority of the similarity measure can be revealed by comparing the probability of creating an error edge. The proposed similarity measure will be introduced in Section 2.2, and its further design in matrix form will be presented in Section 2.3.

2.1 Preliminary

Chernoff Bounds. Sums of random variables have been studied for a long time [4, 10, 25]. Chernoff Bounds [10] provide sharply decreasing bounds that are in the exponential form on tail distributions of sums of independent random variables. It is a widely used fundamental tool in mathematics and computer science [30, 39]. An often used and looser form of Chernoff Bounds is as follows.

Let $X = \sum_1^n X_i$*, where* $X_i = 1$ *with the probability* p_i *and* $X_i = 0$ *with the probability* $1 - p_i$*. All* X_i *are independent random variables. Let* $\mu = \mathbb{E}(X) = \sum_1^n p_i$*. Then for all* $0 < \epsilon < 1$*,*

(i) **Upper Tail:** $\mathbb{P}(X \geq (1+\epsilon)\mu) \leq e^{-\frac{\epsilon^2}{3}\mu}$; (ii) **Lower Tail:** $\mathbb{P}(X \leq (1-\epsilon)\mu) \leq e^{-\frac{\epsilon^2}{3}\mu}$.

It shows that with a high probability $1-\delta$, we have $X \leq (1+\epsilon)\mu$, or $X \geq (1-\epsilon)\mu$, where $\delta = e^{-\frac{\epsilon^2}{3}}$.

GCNs. A GCN network normally consists of multiple GCN layers. Given an input graph $\mathcal{G}(\mathbf{X}, \mathbf{A})$ with L vertices, the output of a normal GCN layer is:

$$\mathbf{X}^{'} = \sigma(\tilde{\mathbf{D}}^{-1}\tilde{\mathbf{A}}\mathbf{X}\mathbf{W}_l), \quad \text{where } \tilde{\mathbf{A}} = \mathbf{A} + \mathbf{I},\ \tilde{\mathbf{D}}_{ii} = \sum_{j=1}^{L} \tilde{\mathbf{A}}_{i,j}, \tag{1}$$

$\mathbf{X} \in \mathbb{R}^{L\times M}$ is input vertex features with M dimensions, $\mathbf{A} \in \mathbb{R}^{L\times L}$ is their adjacency matrix; $\mathbf{I} \in \mathbb{R}^{L\times L}$ is an identity matrix, $\tilde{\mathbf{D}} \in \mathbb{R}^{L\times L}$ is the diagonal degree matrix of $\tilde{\mathbf{A}}$; $\mathbf{W}_l \in \mathbb{R}^{M\times M'}$ is a trainable matrix, and $\sigma(\cdot)$ is an activation function such as ReLU [18]; $\mathbf{X}^{'} \in \mathbb{R}^{L\times M'}$ is the output vertex features with M' dimensions, $M' = M$ in this work.

2.2 Multiple tests on similarity measure

Let $\mathcal{V} = \{v_1, v_2, ..., v_N\}$ be the collection of N nodes. Each node is assumed to be in one of two categories: $+1$ or -1 for simplicity. For each node v_i, the subset of candidate nodes to connect it is denoted by $\mathcal{V}_i$. This subset can be decided by a simple criterion such as by k nearest neighbours. When calculating the proposed similarity between v_i and $v_j \in \mathcal{V}_i$, the common candidate nodes between v_i and v_j, i.e., $\mathcal{V}_{i,j} = \mathcal{V}_i \cap \mathcal{V}_j$ are first identified. The nodes in the $\mathcal{V}_{i,j}$ are treated as the **multiple samples** drawn for the comparison of v_i and v_j. Then, each $v_k \in \mathcal{V}_{i,j}$ performs single test twice, one for v_k and v_i and one for v_k and v_j. The **single test** here is a comparison using the pairwise similarity based on the features of v_k and v_i, or v_k and v_j. The pairwise similarity can be cosine similarity, Gaussian kernel similarity or any other form, and is denoted by **Sim-S**. The proposed similarity score between v_i and v_j increases by one if the two tests yield $C(v_i) = C(v_k)$ and $C(v_k) = C(v_j)$, and zero otherwise, where $C(v)$ is the category of node v. It contains multiple single tests, hence the name **multiple tests**, corresponding similarity measure denoted by **Sim-M**.

For the convenience of analysis, a few assumptions are further introduced. First, the size of $\mathcal{V}_{i,j}$ is m, and among the nodes in $\mathcal{V}_{i,j}$, αm nodes share the same category as v_i and $(1-\alpha)\,m$ nodes belong to the opposite category, with $\alpha > \frac{1}{2}$. For any node $v_j \in \mathcal{V}_i$, its probability of being connected with v_i, according to the statistical test, is $q \in [0, 1]$ if $C(v_i) \neq C(v_j)$, and the probability increases to $1 \geq p > q$ if $C(v_i) = C(v_j)$. This section will show that, if m is sufficiently large (i.e., enough number of tests can be run), there will exist an appropriate threshold for the Sim-M such that the number of error edges can be reduced significantly compared to Sim-S.

Proposition 1. *For the single test method, in expectation, the number of noisy edges is* $(1-\alpha)\,qm$ *and the number of missing edges is* $\alpha\,(1-p)\,m$.

The theorem below shows that both the noisy and missing edges can be reduced by a significant portion when m is not too small.

Theorem 1. *Assume for* $\gamma > 1$*, we have*

$$m > \frac{3((p+q)^2 + (2\alpha-1)(p^2-q^2))^2}{pq((p-q)^2 + (2\alpha-1)(p^2-q^2))^2} \log(\frac{\gamma}{\min((1-\alpha)q, \alpha(1-p))}), \tag{2}$$

Then, for the multiple test method, in expectation, it can reduce the number of noisy edges and missing edges of the single test method by $1/\gamma$.

Proof. Consider node v_i and one of its candidate node $v_j \in \mathcal{V}_i$. $S^k_{i,j}$ is denoted as the similarity score between v_i and v_j based on the statistical test with respect to v_k. As shown in Figure 1, v_k may be of the same category as v_i or not (denoted by v'_k for a better show). Evidently, $\mathbb{E}[S^k_{i,j}] = pq$ if $C(v_i) \neq C(v_j)$, and $\mathbb{E}[S^k_{i,j}] = \alpha p^2 + (1-\alpha)q^2$ if $C(v_i) = C(v_j)$.

The Sim-M between v_i and v_j is defined as:

$$S_{i,j} = \frac{1}{m} \sum_{k\in\mathcal{V}_{i,j}} S^k_{i,j}. \tag{3}$$

Using the Chernoff Bounds [10], we have with a probability $1-\delta$,

$$\begin{cases} S_{i,j} = \frac{1}{m}\sum_{k\in V_{i,j}} S_k \leq (1+\sqrt{\frac{3log(1/\delta)}{pqm}})pq, & C(v_i) \neq C(v_j), \\ S_{i,j} = \frac{1}{m}\sum_{k\in V_{i,j}} S_k \geq (1-\sqrt{\frac{3log(1/\delta)}{(\alpha p^2+(1-\alpha)q^2)m}})(\alpha p^2 + (1-\alpha)q^2), & C(v_i) = C(v_j), \end{cases} \tag{4}$$

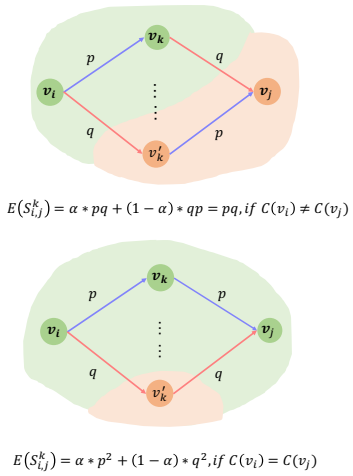


Figure 1: Illustration of $\mathbb{E}[S_{i,j}^k]$ in Sim-M on two nodes from different categories and the same category. Nodes of different colors are from different categories.

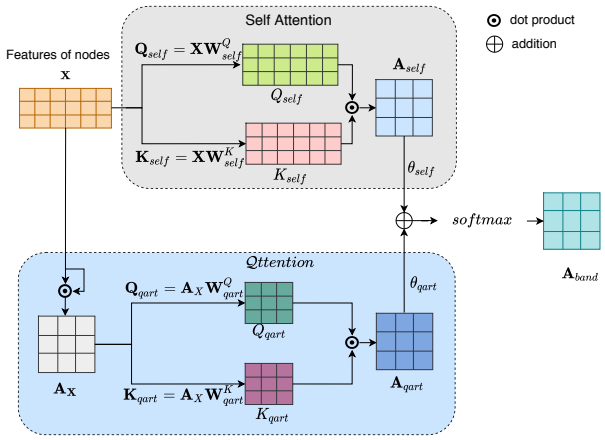


Figure 2: Illustration of the $\mathcal{B}$**ttention** mechanism. The self-attention part is the same as that in Transformers. The $\mathcal{Q}$**ttention** part generates $\mathbf{A}_X$ and then pay attention to it to generate the $\mathbf{A}_{qart}$. The two output attention maps are fused as the final output $\mathbf{A}_{band}$.

By assuming that m is large enough, i.e.,

$$\sqrt{\frac{3\log(1/\delta)}{pqm}} < \frac{(p-q)^2 + (2\alpha-1)(p^2-q^2)}{(p+q)^2 + (2\alpha-1)(p^2-q^2)}, \tag{5}$$

by choosing the similarity threshold

$$S_t = (1 + \sqrt{\frac{3log(1/\delta)}{pqm}})\frac{pq}{2} + (1 - \sqrt{\frac{3log(1/\delta)}{(\alpha p^2 + (1-\alpha)q^2)m}})\frac{\alpha p^2 + (1-\alpha)q^2}{2}, \tag{6}$$

only with a probability δ, we will miss a correct edge, and with a probability δ we will introduce a noisy edge. We complete the proof by substituting δ with

$$\frac{1}{\gamma}\min((1-\alpha)q, \alpha(1-p)). \tag{7}$$

We can guarantee that the Sim-M between two nodes from the same category is strictly larger than that for two nodes from different categories, leading to the distillation of all noisy edges.

2.3 Multiple tests on GCNs

With a better similarity measure prototype above, we further craft it into an elegant and efficient matrix form for adaptation of graph structure learning in the real-world. Given the L2 normalized original node features $\mathbf{X}$, the idea of Sim-S can be implemented by inner product of features, i.e. $\mathbf{A}_X = \mathbf{X}\mathbf{X}^T$. The twice-run tests on each node can be achieved by $\mathbf{A}_X$ multiplying itself. Inspired by the design of popular Transformers [50], some learnable parameters are also introduced. It contains the **fourth-order statistics of features**, which is a significant difference, since self-attention only has second-order. If self-attention is regarded as **Duet of Attention**, our method is analogous to **Quertet of Attention** ($\mathcal{Q}$**ttention** for short). The vectorized representation of Equation 3, $\mathcal{Q}$**ttention** $\mathbf{A}_{qart} \in \mathbb{R}^{L\times L}$, is then implemented as follows:

$$\mathbf{A}_{qart} = \mathbf{Q}_{qart}\mathbf{K}_{qart}^T, \quad \text{where } \mathbf{Q}_{qart} = \mathbf{A}_X\mathbf{W}_{qart}^Q,\ \mathbf{K}_{qart} = \mathbf{A}_X\mathbf{W}_{qart}^K, \tag{8}$$

$\mathbf{W}_{qart}^Q \in \mathbb{R}^{L\times L'}$ and $\mathbf{W}_{qart}^K \in \mathbb{R}^{L\times L'}$ are learnable weights, $L' = L$ in this work.

Motivated by the further enhancements from combinations of intrinsic graph structures and implicit graph structures in literature [33, 7, 34], the self-attention can play an intrinsic role to fuse with the

$\mathcal{Q}$**ttention** in vision tasks, although no intrinsic graph structure exists between images at all. The **self-attention** part is the same as that in Transformers [50]. The attention map $\mathbf{A}_{self} \in \mathbb{R}^{L\times L}$ is:

$$\mathbf{A}_{self} = \frac{\mathbf{Q}_{self}{\mathbf{K}_{self}}^T}{\sqrt{M^d}}, \quad \text{where } \mathbf{Q}_{self} = \mathbf{X}\mathbf{W}^Q_{self},\ \mathbf{K}_{self} = \mathbf{X}\mathbf{W}^K_{self}, \tag{9}$$

$\mathbf{W}^Q_{self} \in \mathbb{R}^{M\times M^d}$ and $\mathbf{W}^K_{self} \in \mathbb{R}^{M\times M^d}$ are the learnable weights, M^d is the dimension of query and key features. The combined form blends the duet and quartet of attention, hence the name **Band of Attention** ($\mathcal{B}$**ttention** for short). The overview of $\mathcal{B}$**ttention** mechanism is shown in Figure 2.

The fusion of $\mathbf{A}_{self}$ and $\mathbf{A}_{qart}$ is defined as (where softmax$(\cdot)$ is a softmax function):

$$\mathbf{A}_{band} = \text{softmax}(\theta_{qart}\mathbf{A}_{qart} + \theta_{self}\mathbf{A}_{self}). \tag{10}$$

When combined with the neural network layer in Equation 1, the final form is:

$$\mathbf{X}' = \sigma(\mathbf{A}_{band}\mathbf{X}\mathbf{W}_l). \tag{11}$$

In real implementation, considering the graph size can be very large for large-scale datasets, sub-graph sampling strategy [53, 52] is adopted to make it more computationally efficient and scalable. In particular, k_{seed} nodes are first selected as seeds from a dataset in each sampling step. Then the seed nodes together with their k nearest neighbours (kNN) [13] form the nodes of a sub-graph. As a result, Equations 8 and 9 are applied to the sub-graph during training and inference. For a dataset of size N_d, it takes about N_d/k_{seed} iterations to process the whole dataset on average.

3 Experiments

3.1 Experimental setups

Datasets Experiments are conducted on commonly used visual datasets[1] with three different types of objects, i.e., **MS-Celeb** [20] (human faces), **MSMT17** [55] (human bodies) and **VeRi-776** [35] (vehicles), to verify the generalization of the proposed method. In these datasets, each sample has a specific category, and most categories have rich samples. The tasks on these datasets are popular and representative in computer vision. It is worth noting that MS-Celeb is divided into 10 parts by identities (Part0-9): Part0 for training and Part1-9 for testing, to maintain comparability and consistency with previous works [59, 59, 19, 53, 40, 47]. **Details of the dataset** are in Section A.1.

Metrics The Area Under the Curve (**AUC**) is adopted to directly evaluate the discriminatory power of similarity measures between node pairs. Edge Noise Rate (**ENR**) illustrates the amount of noise in graphs quantitatively. For node v_i, $\mathbf{ENR} = \frac{N_i^{noise}}{N_i^{all}}$, where N_i^{noise} is the number of noisy edges and N_i^{all} is the amount of all edges for v_i. The average ENR is defined as the average of the ENRs of all nodes. The feature quality of GCNs output is investigated to prove the effectiveness of graph structure learning: The mean Average Precision (**mAP**) [63] is used from the perspective of identification and **ROC** curves are shown from the perspective of verification. The higher the quality of the features the better the graph structure learning. In downstream ReID tasks, $\mathbf{mAP^R}$ and Rank1 accuracy (**R1**) are used. Compared to mAP, mAP$^{\mathbf{R}}$ additionally takes into account the camera id information, i.e., samples from the same camera as the probe will not be counted in evaluation. **BCubed** ($\mathbf{F_B}$) [3, 1] and **Pairwise** ($\mathbf{F_P}$) [48] F-scores are used for evaluating clustering tasks.

3.2 Effectiveness of the Sim-M

To verify the robustness of Sim-M, the study is first conducted on MS-Celeb (Part1) by evaluating the discriminatory ability on different average ENRs comparing with Sim-S. The Sim-S uses the cosine similarity here for simplicity and without loss of generality. For the sake of sparsity and efficiency it brings, the graphs are built based on kNN via the approximate nearest neighbour algorithm. The average ENR of graphs is controlled by using different k values when searching neighbours. AUC

[1] **DECLARATION:** Considering the ethical and privacy issues, all the datasets used in this paper are feature vectors which can not be restored to images.

is adopted to evaluate the similarity score between node pairs which have kNN relations. Table 1 presents that as the k increases, more and more noise is added (ENR gets larger). For Sim-M, however, more samples can be drawn and more tests can be run, so the advantage of Sim-M over Sim-S becomes increasingly obvious. This demonstrates the **robustness** of Sim-M to noise. This phenomenon also conforms to the condition and conclusion in **Theorem 1**. However, as k increases and ENR gets larger, too much noise can not be handled by multiple tests, so the performance of Sim-M begins to drop. This downward trend can be slowed down by the careful design of $\mathcal{B}$**ttention**. The same experiments are conducted on the MSMT17 and VeRi-776, with similar findings in Section A.2.

3.3 Effectiveness of the $\mathcal{B}$ttention

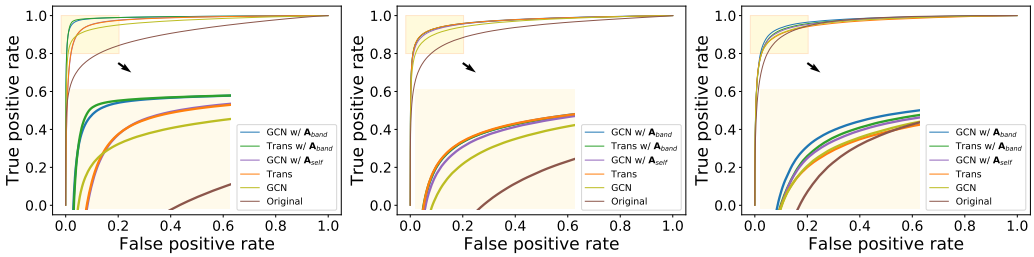
To demonstrate the robustness of $\mathcal{B}$**ttention** mechanism, the study is first conducted on MS-Celeb (Part1) by evaluating the performance of $\mathcal{B}$**ttention** with different average ENRs, comparing it to other baselines. Then, comparison experiments are also conducted on MSMT17, VeRi-776 and also larger scales (Part3-9) of MS-Celeb with the best ENR settings for each dataset. Comparison baselines include the original features of each dataset, a commonly used form of GCN [32, 53], Transformer, GCN with the self-attention (GCN with $\mathbf{A}_{self}$), Transformer with the $\mathcal{B}$**ttention** mechanism (Trans with $\mathbf{A}_{band}$), and GCN with $\mathcal{B}$**ttention** mechanism (GCN with $\mathbf{A}_{band}$). Transformer here is only used as an encoder for structure learning, no positional encoding is required.

Table 1: Average ENRs of different k values and the AUC of Sim-S ($\text{AUC}_\mathbf{S}$), Sim-M ($\text{AUC}_\mathbf{M}$) and their difference (AUC_δ=$\text{AUC}_\mathbf{M}$-$\text{AUC}_\mathbf{S}$) on MS-Celeb (part1). The values of AUC are scaled by a factor of 100 for better display.

k	ENR	$\text{AUC}_\mathbf{S}$	$\text{AUC}_\mathbf{M}$	AUC_δ
5	0.05	96.80	97.59	0.79
10	0.09	96.21	97.72	1.51
20	0.14	95.33	97.21	1.88
40	0.23	94.00	95.66	1.66
80	0.39	92.05	91.32	-0.71
120	0.54	90.76	86.58	-4.17
160	0.64	90.65	83.90	-6.75

Network architectures and settings The model is composed of multiple GCN layers with $\mathcal{B}$**ttention**, and an extra fully connected layer with PReLU [22] activation. The output features have a dimension of 2048 for all datasets. The number of GCN layers is also tuned respectively for different datasets: MS-Celeb uses three GCN layers; MSMT17 and VeRi-776 use one GCN layer. Each node acts as the probe to search its kNN to construct the graph. The k value is tuned respectively for different datasets: 120 for MS-Celeb, 30 for MSMT17, and 80 for VeRi-776. Each node in a dataset can be enhanced multiple times, as either a probe or a neighbour. The Hinge Loss [43, 53] is used to classify node pairs. The initial learning rate is 0.008 with the cosine annealing strategy.

Performance on different average ENRs Models are trained on Part0 and tested on Part1 of MS-Celeb. All baselines use three GCN/Transformer layers for MS-Celeb, except the GCN with $\mathbf{A}_{self}$ which uses two GCN layers (observed to have better performance than three GCN layers). For a certain ENR, the same k value is used in training and testing. Figure 3 shows how the output feature qualities (mAP) change with different average ENRs for all baselines. It shows that the GCN and Transformer with $\mathbf{A}_{band}$ work the best in most average ENRs in terms of both mAP values and their robustness to different ENRs. The GCN has the worst results, and the performances of the ones with $\mathbf{A}_{self}$ are in between. This validates that a commonly used form of GCN is sensitive to noisy edges in input graphs; the proposed $\mathcal{B}$**ttention** can well alleviate the influence of noisy edges and show the strongest robustness when learning graph structure; the $\mathbf{A}_{self}$ also has some power of dealing with noisy edges, but it is worse than $\mathbf{A}_{band}$. When k value is too small (e.g., 5, 10), the performances of $\mathbf{A}_{band}$ and $\mathbf{A}_{self}$ are similar. This indicates that the $\mathcal{Q}$**ttention** part of $\mathcal{B}$**ttention** can hardly bring a positive effect when there are too few context neighbours, i.e. the number of single tests gathered is insufficient. In addition, GCN and Transformer have very similar performance, either with $\mathbf{A}_{band}$ or $\mathbf{A}_{self}$. This indicates that base networks are not critical, and both attention mechanisms work for the two base networks. The performance degradation of $\mathcal{B}$**ttention** with very large ENR (e.g., 0.64) indicates that the current form $\mathcal{B}$**ttention** still has improvement room, although works much better than the self-attention.



(a) MS-Celeb Part1 (b) MSMT17 (c) VeRi-776

Figure 4: ROC curves of different methods on MS-Celeb Part1 (a), MSMT17 (b) and VeRi-776 (c).

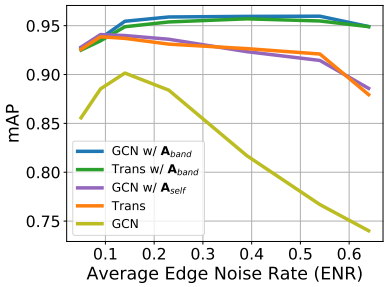


Figure 3: Feature quality (mAP) of various methods with different average ENRs on MS-Celeb Part1.

Table 2: Feature quality (mAP) of different methods on visual datasets of different types of objects.

Dataset	MS-Celeb	MSMT17	VeRi-776
Original Feature	80.07	74.46	81.91
GCN	90.15	83.51	85.27
Transformer	93.86	85.44	85.92
GCN w/ $\mathbf{A}_{self}$	94.09	85.40	86.88
Trans w/ $\mathbf{A}_{band}$	95.70	85.92	87.07
GCN w/ $\mathbf{A}_{band}$	**95.97**	**86.04**	**87.59**

Performance on different types of objects To further validate the $\mathcal{B}$**ttention** mechanism, comparisons are also made on datasets with other types of objects: MSMT17 and VeRi-776. In these two datasets, k values and network depths are also **tuned** for all baselines, aiming for their best performances. For both datasets, one GCN layer is used for GCN, GCN with $\mathbf{A}_{band}$, and Transformer with $\mathbf{A}_{band}$; three GCN layers are used for GCN or Transformer with $\mathbf{A}_{self}$. In MSMT17, k=10 for GCN and "GCN w/ $\mathbf{A}_{self}$"; k=20 for Transformer; k=30 for the two $\mathcal{B}$**ttention** methods. In VeRi-776, k=80 for "GCN w/ $\mathbf{A}_{band}$" and "Trans w/ $\mathbf{A}_{band}$", and k=20 for the rest of baselines. The best mAP results of all baselines are listed in Table 2, where the results for MS-Celeb are the peak values in Figure 3. From Table 2, the $\mathcal{B}$**ttention** works the best in all three datasets; $\mathbf{A}_{self}$ works better than GCN; and all output features are much better than the original ones, although the original features of MSMT17 and VeRi-776 are current SOTA ones. The best performances are achieved by the GCN with $\mathbf{A}_{band}$ in all datasets. Their ROC curves are shown in Figure 4, where the same observations can be seen from verification perspective. As shown in Figure 4 and Table 2, the performance differences among those baselines with attention mechanisms are less significant in MSMT17 where k values are relatively small and close across baselines. This observation is similar to that in Figure 3: the performance margin between $\mathbf{A}_{band}$ and $\mathbf{A}_{self}$ is less significant when average ENRs (i.e., k) are relatively small. This further shows that the benefit of $\mathcal{B}$**ttention** is limited when with a small number of neighbours, as not many single tests to run and leverage. Better performance can not be achieved by enlarging k values for MSMT17, as its average sample amount for each individual is only 31. Too large k values would result in too big ENR which can not be handled by the current form of $\mathcal{B}$**ttention** as discussed on Figure 3.

Experiments are also conducted to compare these baselines on **larger scales of MS-Celeb** (i.e., Part3-9) in Section A.3 of Appendix. Results in Table 6 and Figure 5 show that all baselines have performance degradation when the scale enlarges while the degradation of that "w/ $\mathbf{A}_{band}$" is much smaller than baselines. These results for the cases with different ENRs, on different types of objects and larger scales of testsets, together validate the robustness and effectiveness of proposed $\mathcal{B}$**ttention** in graph structure learning.

Ablation experiments on the design of the topological architecture and attention fusion method are also conducted in Section A.4 of Appendix.

Table 3: Clustering performance on MS-Celeb.

Dataset	Part1		Part3		Part5		Part7		Part9	
Metrics	F_P	F_B	F_P	F_B	F_P	F_B	F_P	F_B	F_P	F_B
K-Means	79.21	81.23	73.04	75.20	69.83	72.34	67.90	70.57	66.47	69.42
HAC	70.63	70.46	54.40	69.53	11.08	68.62	1.40	67.69	0.37	66.96
DBSCAN	67.93	67.17	63.41	66.53	52.50	66.26	45.24	44.87	44.94	44.74
Clusformer	88.20	87.17	84.60	84.05	82.79	82.30	81.03	80.51	79.91	79.95
STAR-FC	91.97	90.21	88.28	86.26	86.17	84.13	84.70	82.63	83.46	81.47
Ada-NETS	92.79	91.40	89.33	87.98	87.50	86.03	85.40	84.48	83.99	83.28
Original+G-cut	69.63	73.62	63.31	66.56	60.61	62.91	57.13	58.72	54.42	58.54
Ours+G-cut	**93.90**	**92.47**	**90.54**	**89.23**	**88.70**	**87.13**	**86.18**	**85.59**	**84.36**	**84.10**
Original+Infomap	93.91	92.42	90.10	89.08	88.49	87.30	85.31	86.01	83.08	85.01
Ours+Infomap	**94.94**	**93.67**	**91.74**	**90.81**	**89.50**	**89.15**	**87.04**	**87.81**	**85.40**	**86.76**

3.4 Effectiveness of the downstream tasks

The improvement of graph structure learning can also be proven through the downstream tasks by exploiting the features it produces. The better the graph structure is learned, the better the performance on downstream tasks. Graphs have important applications in large-scale clustering and re-reranking in ReID. However, they suffer from inaccurate weight problems when learning and working with the graph structure. Armed the GCN with $\mathbf{A}_{band}$, many SOTA results are achieved.

Performance on clustering tasks The clustering tasks are evaluated on MS-Celeb and MSMT17. Both of them have a large number of classes and are suitable as clustering benchmarks. The same features output by "GCN w/ $\mathbf{A}_{band}$" are used as that in Section A.3 for MS-Celeb and Section 3.3 for MSMT17. The same clustering method (denoted "G-cut") is adopted as that in [19, 53], where pairs of nodes are linked when their similarity scores are larger than a tuned threshold, and clustering is finally completed by transitively merging all links via a union-find algorithm [2]. The threshold is tuned for each dataset, aiming for the best balance between BCubed and Pairwise F-Scores. In addition, map equation based unsupervised clustering algorithm Infomap [44, 57] is also adopted here. Their BCubed and Pairwise F-scores are listed in Table 3 and Table 10, where the results of current SOTA GCN/Transformer methods (Clusformer [40], STAR-FC [47] and Ada-NETS [53]) and three conventional clustering approaches (K-Means [36], HAC [49] and DBSCAN [16]) are also listed. The results show that the features enhanced via our approach have the best clustering results in all datasets, either with G-cut or Infomap, both of which outperform the current SOTA methods Ada-NETS and STAR-FC with large margins. In comparison, the results of the ones with Infomap are better than those with G-cut.

Table 4: ReID performance on MSMT17 and VeRi-776.

Datasets	MSMT17		VeRi-776	
Metrics	mAP^R	R1	mAP^R	R1
Original Feature	62.76	82.36	79.00	96.60
GCN	74.66	85.36	82.31	96.60
Transformer	77.60	86.37	82.62	96.54
GCN w/ $\mathbf{A}_{self}$	77.34	85.49	84.29	96.30
Trans w/ $\mathbf{A}_{band}$	77.49	86.54	83.98	96.72
GCN w/ $\mathbf{A}_{band}$	78.28	**86.64**	**84.74**	96.78
Original+QE	71.10	84.37	83.09	95.47
Original+LBR	70.36	85.28	83.63	**97.13**
Original+KR	76.46	85.48	82.32	96.84
Original+ECN	**78.79**	86.16	84.25	**97.13**

Performance on ReID tasks The ReID tasks is evaluated on two ReID datasets: MSMT17 and VeRi-776. The enhanced features used are the same as those in Table 2. mAP$^{\mathbf{R}}$ and R1 are evaluated following the standard probe and gallery protocols of MSMT17 and VeRi-776. The results are shown in Table 4. As the output feature of learned graph can also be understood as a rerank process in ReID, comparisons are also made on commonly used rerank methods: QE [12], LBR [37], KR [62] and ECN [45]. Table 4 shows that the "GCN w/ $\mathbf{A}_{band}$" achieves the best or comparable performance, and all enhanced features are much better than the original SOTA ones. In MSMT17, similar to the results in Section 3.3, the baselines with attention mechanism have very close performance, but work much

better than the GCN. In VeRi-776, from the mAP$^{\mathbf{R}}$ perspective, $\mathcal{B}$**ttention** works better than $\mathbf{A}_{self}$, and both of them are better than the GCN, but their R1 results are comparable. These results still support the superiority of the proposed approach, although R1 results of VeRi-776 are comparable, as mAP$^{\mathbf{R}}$ is a more comprehensive indication of feature quality for ReID. The experiments on the **combination** of "GCN w/ $\mathbf{A}_{band}$" with these commonly used rerank methods are also conducted in Section A.6, achieving more SOTA performances and proving the complementarity between them.

4 Related Work

Graph Convolutional Networks. GCNs [32, 21, 51] extends the operation of convolution from the grid-data (e.g. images, videos) to non-grid data which is more common in the real world. The classic GCN [32] is proposed based on the spectral theory and achieves promising results on citation network datasets: Citeseer, Cora and Pubmed [46]. GraphSAGE [21] further extends GCNs from transductive framework to inductive framework and obtains better results. GAT [51] introduces the attention mechanism in GCN, making it more expressive. Fast-GCN [6] interprets graph convolutions as integral transforms of embedding functions under probability measures, which not only is efficient for training but also generalizes well for inference. However, these GCNs are all investigated on relational datasets that have explicit graph structures. This work aims to extend the capacity of GCNs to work on datasets (e.g., image objects) which have no explicitly defined graph structures.

Graph Structure Learning. The literature [56] suggests that research on graph structure learning for GCNs can be roughly divided into discrete-based and weighted-based ones. Some discrete-based methods [15, 29, 38, 17] treat the graph structures as random distributions and sample binary adjacency matrices from them. Another way [53] constructs discrete graphs based on heuristic proxy objectiveness aiming for clean yet rich neighbours for each node. The binary matrices methods are difficult to optimize because of the gradient breakage, so such methods resort to variational inference (e.g., approaches in [15, 38]), reinforcement learning (e.g., approaches in [29]) or optimization in stages (e.g., approaches in [53]), which may lead to sub-optimal solutions. In addition, the parameter size of distributions [17] can be very large (e.g., $\mathcal{O}(Nk)$ where N is the number of nodes and k is the number of neighbours for each node) and therefore not suitable for large datasets. The weighted-based methods [8, 9, 41, 60, 26, 7, 24, 33, 19, 40], which are the focus of this paper, learn weighted adjacency matrices so that it can be optimized by SGD techniques [42, 31, 5]. The main difference between these works [8, 9, 41, 60, 26, 7, 24, 33] is in the similarity measure, which can be classified into attention-based [8, 9, 41, 60, 26, 7] and kernel-based [24, 33] roughly. To enhance the discriminatory power of the similarity measure, they have adopted various forms: adding learnable parameters [8, 26, 7], introducing nonlinearity [9, 41, 60], Gaussian kernel functions [24, 33] or self-attention [19, 40, 50, 11] etc. However, these methods are sensitive to the noise from feature representations, particular when each node is represented by a high dimensional vector, leading to the problem of inaccurate weights and vulnerable graph structure. This work aims to propose a multiple samples based similarity measure to improve the robustness of graph structure learning over images. Another approach uses the low-rank, sparsity and feature smoothness properties of the real-world graph to optimize the graph structure [27], but these priors are suitable for community networks rather than graph of image objects. Some work [28] learns to augment the given graph with new edges by reinforcement learning to improve the performance of downstream tasks, but existing noisy edges problem in graphs cannot be handled.

5 Conclusion

The problem of inaccurate edge weights is identified in graph structure learning, analysed from the perspective of noise in feature vectors, and handled based on several tests. A novel and robust similarity measure Sim-M is proposed, proven to be superior and verified in experiments. Then it is designed in an elegant matrix form, i.e., $\mathcal{B}$**ttention** mechanism, to improve the graph structure quality in GCNs. The superiority and robustness of $\mathcal{B}$**ttention** are validated by comprehensive experiments and achieve many SOTA results on clustering and ReID benchmarks.

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Checklist

1. For all authors...
   (a) Do the main claims made in the abstract and introduction accurately reflect the paper's contributions and scope? [Yes] See abstract and Section 1.
   (b) Did you describe the limitations of your work? [Yes] See Section B in Appendix.
   (c) Did you discuss any potential negative societal impacts of your work? [Yes] See Section C in Appendix.
   (d) Have you read the ethics review guidelines and ensured that your paper conforms to them? [Yes] We have read the ethics guidelines on the official website.
2. If you are including theoretical results...
   (a) Did you state the full set of assumptions of all theoretical results? [Yes] See Section 2.2.
   (b) Did you include complete proofs of all theoretical results? [Yes] See Section 2.2
3. If you ran experiments...
   (a) Did you include the code, data, and instructions needed to reproduce the main experimental results (either in the supplemental material or as a URL)? [No] The code and the data are proprietary. The instructions is in Section 3.
   (b) Did you specify all the training details (e.g., data splits, hyperparameters, how they were chosen)? [Yes] See in Section 3.1, paragraph of *Network architectures and settings* and the paragraphs describing the corresponding experiment in Section 3.3.
   (c) Did you report error bars (e.g., with respect to the random seed after running experiments multiple times)? [No] Error bars are not reported because it would be too computationally expensive.
   (d) Did you include the total amount of compute and the type of resources used (e.g., type of GPUs, internal cluster, or cloud provider)? [Yes] See Section A.7.
4. If you are using existing assets (e.g., code, data, models) or curating/releasing new assets...
   (a) If your work uses existing assets, did you cite the creators? [Yes] See the *Datasets* paragraph in Section 3.1.
   (b) Did you mention the license of the assets? [No] The used assets are very common in papers of computer vision.
   (c) Did you include any new assets either in the supplemental material or as a URL? [No]
   (d) Did you discuss whether and how consent was obtained from people whose data you're using/curating? [No] The used assets are very common in papers of computer vision.
   (e) Did you discuss whether the data you are using/curating contains personally identifiable information or offensive content? [Yes] See the footnote in *Datasets* paragraph in Section 3.1: Considering the ethical and privacy issues, all the datasets used in this paper are feature vectors which can not be restored to images.
5. If you used crowdsourcing or conducted research with human subjects...
   (a) Did you include the full text of instructions given to participants and screenshots, if applicable? [N/A]

(b) Did you describe any potential participant risks, with links to Institutional Review Board (IRB) approvals, if applicable? [N/A]

(c) Did you include the estimated hourly wage paid to participants and the total amount spent on participant compensation? [N/A]

A Experiments

A.1 Details of the dataset

MS-Celeb [20] is a dataset with 10M face images of nearly 100K individuals. A cleaned version [14] with about 5.8M face images of 86K identities is used in this paper. The protocol and features are the same as that in [59], where the dataset is divided into 10 parts by identities (Part0-9): Part0 for training and Part1-9 for testing. The feature dimension is 256. The data amounts of Part 0-9 (0, 1, 3, 5, 7 and 9) are respectively: 584K, 584K, 1.74M, 2.89M, 4.05M and 5.21M.

MSMT17 [55] is the current largest ReID dataset. It contains 32,621 images of 1,041 individuals for training and 93,820 images of 3,060 individuals for testing. These images are captured by 15 cameras under different time periods, weather and light conditions. This work uses the SOTA features from AGW [61] with a dimension of 2048.

VeRi-776 [35] is a commonly used vehicle ReID benchmark, which contains over 40K bounding boxes of 619 vehicles captured by 20 cameras in unconstrained traffic scenes. The dataset is divided into two parts: 37,715 images of 576 vehicles for training, and 13,257 images of 200 vehicles for testing. This work uses the features extracted by the SOTA ViT-B/16 Baseline in TransReID [23], with 768 dimensions.

A.2 Effectiveness of the Sim-M

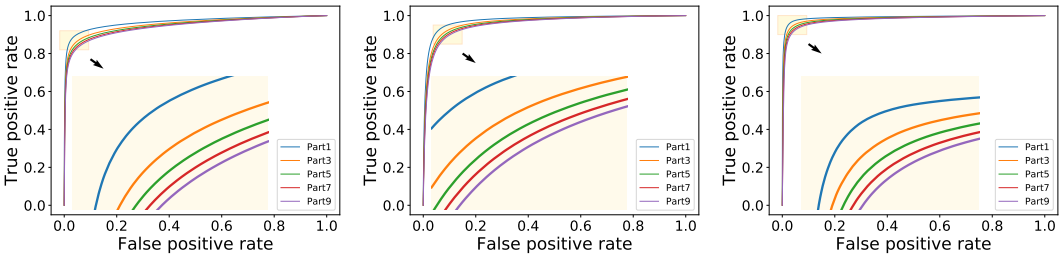
The experiments on MSMT17 and VeRi-776 also show that the advantage of Sim-M over Sim-S is increasingly obvious as k increases and then drops for introduction of too much noise as in Table 5.

Table 5: AUC of Sim-S ($\mathrm{AUC_S}$), Sim-M ($\mathrm{AUC_M}$) and their difference (AUC_δ=$\mathrm{AUC_M}$-$\mathrm{AUC_S}$) on MSMT17 and VeRi-776. The values of AUC are scaled by a factor of 100 for a better show.

Datasets	MSMT17				VeRi-776			
k	ENR	$\mathrm{AUC_S}$	$\mathrm{AUC_M}$	AUC_δ	ENR	$\mathrm{AUC_S}$	$\mathrm{AUC_M}$	AUC_δ
5	0.06	89.23	89.74	0.52	0.01	90.96	90.40	-0.56
10	0.13	89.33	90.02	0.69	0.03	89.34	90.09	0.75
20	0.26	87.78	88.63	0.85	0.07	85.86	86.54	0.68
40	0.48	86.72	86.62	-0.10	0.17	83.05	82.89	-0.16

A.3 Performance on larger scales of MS-Celeb

Experiments are also conducted to compare these baselines on larger scales of MS-Celeb (i.e., Part3-9). The same model for MS-Celeb Part1 is used as in Section 3.3, but tested on Part3-9. The results in Table 6 show that all baselines have performance degradation when the scale enlarges. The $\mathcal{B}$**ttention** obtains the best results in all parts; $\mathbf{A}_{self}$ works better than GCN; all output features are much better than the original ones. Among the baselines with attention mechanisms, the ones using GCN-based networks have better performances. The GCN with $\mathbf{A}_{band}$ has the highest mAP scores. Figure 5 shows the ROC curves of GCN, "GCN w/ $\mathbf{A}_{self}$" and "GCN w/ $\mathbf{A}_{band}$" across all parts of MS-Celeb, from which we can also see the performance degradation with larger testsets. The performance degradation of "GCN w/ $\mathbf{A}_{band}$" is much smaller than the other two baselines, which is shown more clearly from the differences of AUC in Table 7. In Table 7, differences on AUC of two adjacent parts (e.g., part1 - part3, part3 - part5, part5 - part7, part7 - part9) and their summation for each method are shown. It can be observed that the summation of the differences of "GCN w/ $\mathbf{A}_{band}$" is 1.49, which is only 58.30% and 67.26% of the ones of GCN (2.56) and "GCN with $\mathbf{A}_{self}$" (2.22). This validates that the features from "GCN w/ $\mathbf{A}_{band}$" have a much smaller performance degradation than the other two baselines with larger testsets from the perspective of verification. Similar observations can also be seen in Table 6, where the performance degradation of the baselines with attention mechanisms is much smaller than GCN and original features.



(a) GCN (b) GCN with $\mathbf{A}_{self}$ (c) GCN with $\mathbf{A}_{band}$

Figure 5: ROC curves of GCN (a), GCN with $\mathbf{A}_{self}$ (b), and GCN with $\mathbf{A}_{band}$ (c) on MS-Celeb Part1-9.

Table 6: Feature quality (mAP) on different scales of MS-Celeb.

Partition	Part1	Part3	Part5	Part7	Part9
Original Feature	80.07	75.28	73.21	71.73	70.63
GCN	90.15	86.27	84.34	82.95	81.90
Transformer	93.86	91.32	89.99	89.03	88.28
GCN w/ $\mathbf{A}_{self}$	94.09	91.66	90.38	89.45	88.73
Trans w/ $\mathbf{A}_{band}$	95.70	93.38	92.08	91.09	90.30
GCN w/ $\mathbf{A}_{band}$	**95.97**	**93.71**	**92.38**	**91.30**	**90.46**

Table 7: Differences on AUC (δ_{AUC}) of two adjacent parts of MS-Celeb for GCN, "GCN with $\mathbf{A}_{self}$" and "GCN with $\mathbf{A}_{band}$". The values in the table are scaled by a factor of 100 for better display.

δ_{AUC}	GCN	GCN with $\mathbf{A}_{self}$	GCN with $\mathbf{A}_{band}$
part1 - part3	1.23	1.03	0.58
part3 - part5	0.58	0.51	0.37
part5 - part7	0.43	0.37	0.30
part7 - part9	0.32	0.31	0.25
summation	2.56	2.22	1.49

A.4 Ablation study on $\mathcal{B}$ttention

Ablation experiments are conducted mainly on MS-Celeb to study the design of the topological architecture and attention fusion method of $\mathcal{B}$**ttention**. Similar to Section 3.3, models are trained on Part0 and tested on Part1. All settings other than the factor being investigated are the same as that in Section 3.3 for "GCN with $\mathbf{A}_{band}$" with $k = 120$.

A.4.1 topological architecture design

This study compares the topological architecture of the $\mathcal{B}$**ttention** with three other designs: (1) one with only the $\mathbf{A}_{qart}$ of $\mathcal{B}$**ttention**; (2) one using only the $\mathcal{Q}$**ttention** part but with the $\mathbf{A}_X$ replaced by $\mathbf{A}_{self}$ in Equation 8, denoted as $\widetilde{\mathbf{A}}_{qart}$; (3) and one with $\mathcal{B}$**ttention** but with the $\mathbf{A}_{qart}$ replaced by $\widetilde{\mathbf{A}}_{qart}$, denoted as $\widetilde{\mathbf{A}}_{band}$. Table 8 shows their mAP results, where the results of "GCN w/ $\mathbf{A}_{band}$" and one with only $\mathbf{A}_{self}$ are also listed for comparison. For the purpose of ablation study, the one with only $\mathbf{A}_{self}$ here has the same depth (i.e., three layers) with "GCN w/ $\mathbf{A}_{band}$", which is different from that in Table 2 (two layers). The comparisons are made on two conditions: with or without $\mathbf{W}_{qart}$ (i.e., $\mathbf{W}^Q_{qart}$ and $\mathbf{W}^K_{qart}$) in Equation 8. For the cases without $\mathbf{W}_{qart}$, $\mathbf{W}^Q_{qart}$ and $\mathbf{W}^K_{qart}$ are both identity matrices, i.e., $\mathbf{Q}_{qart} = \mathbf{K}_{qart} = \mathbf{A}_X$. Their ROC curves are shown in Figure 6. It can be seen that the $\mathcal{B}$**ttention** design achieves the best results in both mAP and ROC; $\mathbf{A}_{qart}$ has better performance than $\mathbf{A}_{self}$; $\widetilde{\mathbf{A}}_{qart}$ and $\widetilde{\mathbf{A}}_{band}$ are worse than $\mathbf{A}_{qart}$ and $\mathbf{A}_{band}$ respectively. This shows that the combination of $\mathbf{A}_{self}$ and $\mathbf{A}_{qart}$ and the use of $\mathbf{A}_X$ (instead of $\mathbf{A}_{self}$) in $\mathbf{A}_{qart}$

both contribute to the performance improvement. In addition, the baselines with $\mathbf{W}_{qart}$ have better performances (higher mAP scores and better ROC) than those without, indicating that $\mathbf{W}^{Q}_{qart}$ and $\mathbf{W}^{K}_{qart}$ are also critical for better graph structure learning.

Table 8: Feature quality (mAP) of different topological architectures of the attention mechanism.

Arch./Cond.	w/ $\mathbf{W}_{qart}$	w/o $\mathbf{W}_{qart}$
$\mathbf{A}_{self}$	88.10	88.10
$\widetilde{\mathbf{A}}_{qart}$	90.10	89.40
$\mathbf{A}_{qart}$	92.26	88.92
$\widetilde{\mathbf{A}}_{band}$	94.19	89.53
$\mathbf{A}_{band}$	**95.97**	**93.67**

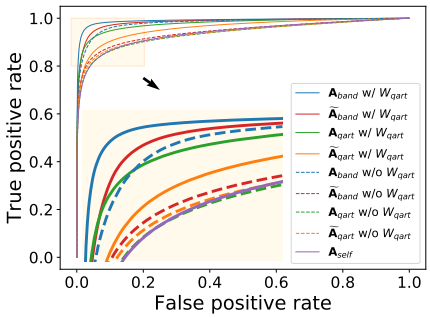


Figure 6: ROC curves of different topological architectures of the attention mechanism.

The ablation experiments are also **conducted on attention map fusion method in Section A.4.2 in Appendix**. The results in Table 9 and Figure 7 together will prove that the proposed form in Equation 11 has the best performance across all datasets.

A.4.2 Attention map fusion method

The attention map fusion method is studied by comparing it to two other designs: direct element-wise addition without weighting ($\mathbf{A}_{qart} + \mathbf{A}_{self}$) and element-wise multiplication ($\mathbf{A}_{qart} \odot \mathbf{A}_{self}$) in Equation 11. The results are shown in Table 9 and Figure 7: The $\theta_{qart}\mathbf{A}_{qart} + \theta_{self}\mathbf{A}_{self}$ design works the best, and the performance of $\mathbf{A}_{qart} \odot \mathbf{A}_{self}$ is very close to $\theta_{qart}\mathbf{A}_{qart} + \theta_{self}\mathbf{A}_{self}$. However, results of MSMT17 and VeRi-776 in Table 9 shows our proposed method has the best performance across all datasets.

Table 9: Feature quality (mAP) of different attention map fusion methods, where $\odot$ denotes element-wise multiplication.

Dataset	$\mathbf{A}_{qart} + \mathbf{A}_{self}$	$\mathbf{A}_{qart} \odot \mathbf{A}_{self}$	$\theta_{qart}\mathbf{A}_{qart} + \theta_{self}\mathbf{A}_{self}$
MS-Celeb	92.83	95.71	**95.97**
MSMT17	84.95	85.59	**86.04**
VeRi-776	85.97	85.53	**87.59**

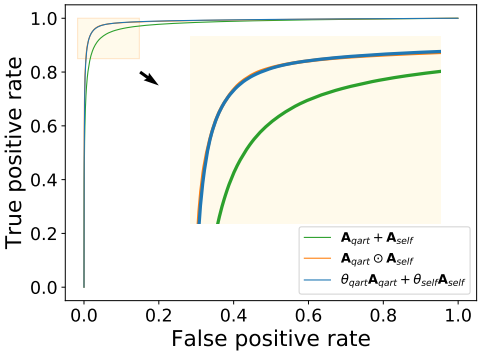


Figure 7: ROC curves of different fusion methods on MS-Celeb (part1).

Table 10: Clustering performance on MSMT17.

Dataset	MSMT17	
Metrics	F_P	F_B
K-Means	60.32	69.28
HAC	63.95	73.75
DBSCAN	48.23	53.90
Clusformer	-	-
STAR-FC	-	-
Ada-NETS	73.51	77.40
Original+G-cut	50.60	66.56
Ours+G-cut	**74.93**	**79.91**
Original+Infomap	71.33	79.16
Ours+Infomap	**77.54**	**81.75**

A.5 Clustering performance on MSMT17

The clustering tasks are also evaluated on MSMT17 and the results are in Table 10.

A.6 Ensemble Performance on ReID tasks

In addition to directly comparing with the commonly used rerank methods (e.g., QE [12], LBR [37], KR [62] and ECN [45]) in Section 4.4, the "GCN with $\mathbf{A}_{band}$" can also be combined with them. The commonly used rerank methods can utilize the enhanced features from "GCN with $\mathbf{A}_{band}$" for better ranking. The results are shown in Table 11. Since mAP$^{\mathbf{R}}$ can provide a more comprehensive evaluation of the feature quality than R1 for ReID, the mAP$^{\mathbf{R}}$ is discussed more often. It can be observed that new SOTA performances are obtained on both datasets. The mAP$^{\mathbf{R}}$ on MSMT17 is as high as 79.67 by "$\mathbf{A}_{band}$+KR" and on VeRi-776 is as high as 85.92 by "$\mathbf{A}_{band}$+LBR". For most of the commonly used rerank methods, the mAP$^{\mathbf{R}}$ of reranking with $\mathbf{A}_{band}$ is higher than with original features and also higher than "GCN w/ $\mathbf{A}_{band}$". The only exception is "$\mathbf{A}_{band}$+ECN" which is higher than "GCN w/ $\mathbf{A}_{band}$" but only comparable with "original+ECN". The results show that there is a complementary relationship between "GCN with $\mathbf{A}_{band}$" and commonly used rerank methods. The "GCN with $\mathbf{A}_{band}$" can provide better features so that the commonly used rerank methods can be further improved, although the $R1$ results are only comparable.

Table 11: Ensemble ReID performance on MSMT17 and VeRi-776. "Original+X" means reranking with original features by methods of "X". "$\mathbf{A}_{band}$+X" means reranking with features from "GCN w/ $\mathbf{A}_{band}$" by methods of "X".

Dataset	MSMT17		VeRi-776	
Metrics	mAP^R	R1	mAP^R	R1
Original Feature	62.76	82.36	79.00	96.60
GCN w/ $\mathbf{A}_{band}$	78.28	86.64	84.74	96.78
Original+QE	71.10	84.37	83.09	95.47
Original+LBR	70.36	85.28	83.63	**97.13**
Original+KR	76.46	85.48	82.32	96.84
Original+ECN	78.79	86.16	84.25	**97.13**
$\mathbf{A}_{band}$+QE	79.37	**86.91**	85.88	96.90
$\mathbf{A}_{band}$+LBR	79.54	86.86	**85.92**	96.90
$\mathbf{A}_{band}$+KR	**79.67**	84.81	85.45	96.60
$\mathbf{A}_{band}$+ECN	78.70	84.05	85.61	96.90

A.7 Declaration of the type of computing resources

All experiments are conducted on a single machine, with 54 E5-2682 v4 CPUs, 8 NVIDIA P100 cards and 400G memory.

B Limitations

Our approach achieves promising results in graph structure learning, but it still has the following limitations:

- The $\mathcal{B}$**ttention** requires more computation and number of parameters as shown in Section 2.3. However, the baseline methods can not achieve the same performance even adding their network depth. The layer number of network in each method is tuned in all experiments.
- The superiority of $\mathcal{B}$**ttention** is not obvious enough on datasets with few samples in each category compared with self attention as revealed in Section 3.3.

- The performance of $\mathcal{B}$**ttention** will also degrade when with large ENR as discussed in Section 3.3, indicating the current form still has improvement room on the robustness against noise to further investigate.
- The outputs $\mathcal{B}$**ttention** will vary depending on the choice of nearest neighbours. However, the superiority of this method can be guaranteed as long as the conditions in Section 2.2 are met.

C Potential Negative Societal Impacts

The proposed graph structure learning method inherently is harmless just like many other AI technologies. This technology may have negative societal impacts if someone uses it for malicious applications such as surveillance, personal information collections. As authors, we advocate for proper technology usage.

D Pseudocode for self-attention and $\mathcal{Q}$ttention

Algorithm 1 PyTorch-style pseudocode for self-attention and $\mathcal{Q}$**ttention**

```
# X: the input feature matrix, with shape (N, D)
def self_attention(X):
    Q_self = torch.matmul(X, W_self_Q)
    K_self = torch.matmul(X, W_self_K)
    self_attn = torch.matmul(Q_self, K_self.transpose(-2, -1))
    return self_attn

def Q_attention(X):
    A_X = torch.matmul(X, X.transpose(-2, -1))
    Q_qart = torch.matmul(A_X, W_qart_Q)
    K_qart = torch.matmul(A_X, W_qart_K)
    Q_attn = torch.matmul(Q_qart, K_qart.transpose(-2, -1))
    return Q_attn
```

E Detailed Proof of with generalized version of Chernoff Bounds

Below is a full analysis for the practical algorithm in Section 2.3 that does not assume a binary relationship between two nodes in A_X.

Let X_i denote a random variable with no distribution assumptions such that $a \leq X_i \leq b$ for all i. Define $X = \sum_1^m X_i$, $\mu = \mathbb{E}(X)$. Then for all $0 < \epsilon < 1$, we have the general version of Chernoff inequality as follows.

(i) **Upper Tail:** $\mathbb{P}(X \geq (1+\epsilon)\mu) \leq e^{-\frac{\epsilon^2\mu^2}{m(b-a)^2}}$;

(ii) **Lower Tail:** $\mathbb{P}(X \leq (1-\epsilon)\mu) \leq e^{-\frac{\epsilon^2\mu^2}{m(b-a)^2}}$.

The above show that with probability at least $1-\delta$, $(1-\epsilon)\mu \leq X \leq (1+\epsilon)\mu$ where $\delta := e^{-\frac{\epsilon^2\mu^2}{m(b-a)^2}}$ or equivalently $\epsilon = \sqrt{\frac{m(b-a)^2 log(1/\delta)}{\mu^2}}$.

Instead of assuming p and q for making connections in the case of $C(v_i) = C(v_j)$ and $C(v_i) \neq C(v_j)$, we assume that the similarity score $s_{i,j}$ is sampled from a bounded distribution (bounded between -1 and +1) with mean of s_+ when $C(v_i) = C(v_j)$, and $s_{i,j}$ is sampled from a bounded distribution (bound between -1 and +1) with mean of s_- when $C(v_i) = C(v_j)$. We of course assume $s_+ > s_-$. In expectation, the number of edges connected under $S^k_{i,j}$ on $\mathcal{V}_{i,j}$ is

$$\begin{cases} \mu^- = s_+ s_- m, & C(v_i) \neq C(v_j) \ , \\ \mu^+ = (\alpha s_+^2 + (1-\alpha) s_-^2) m, & C(v_i) = C(v_j) \ , \end{cases} \tag{12}$$

where $C(v)$ is the category of node v.

It should be noted that $(b-a)^2 = 4$ under our assumption. Therefore, when v_i, v_j are sampled from different categories, we have, with probability at least $1-\delta$,

$$\begin{aligned} \sum_{k \in V_{i,j}} S_{i,j}^k &\leq (1+\epsilon^-)\mu^- \\ \sum_{k \in V_{i,j}} S_{i,j}^k &\leq \left(1 + \sqrt{\frac{4mlog(1/\delta)}{(\mu^-)^2}}\right) \mu^- \\ S_{i,j} = \frac{1}{m} \sum_{k \in V_{i,j}} S_{i,j}^k &\leq \left(1 + \sqrt{\frac{4nlog(1/\delta)}{(s_+ s_- m)^2}}\right) s_+ s_- \ . \end{aligned} \tag{13}$$

When v_i, v_j are sampled from the same category, we have, with probability at least $1-\delta$,

$$\begin{aligned} \sum_{k \in V_{i,j}} S_{i,j}^k &\geq (1-\epsilon^+)\mu^+ \\ \sum_{k \in V_{i,j}} S_{i,j}^k &\geq \left(1 - \sqrt{\frac{4mlog(1/\delta)}{\mu^+}}\right) \mu^+ \\ S_{i,j} = \frac{1}{m} \sum_{k \in V_{i,j}} S_{i,j}^k &\geq \left(1 - \sqrt{\frac{4nlog(1/\delta)}{(\alpha s_+^2 + (1-\alpha) s_-^2)^2 m^2}}\right) (\alpha s_+^2 + (1-\alpha) s_-^2) \ . \end{aligned} \tag{14}$$

The two distributions can be well separated if **the above two bounds do not overlap**. That is, we require that

$$(1+\epsilon^-)\mu^- < (1-\epsilon^+)\mu^+ \ . \tag{15}$$

On the other hand, it is obvious $s_+ s_- < \alpha s_+^2 + (1-\alpha) s_-^2$, so we know

$$\epsilon^- = \sqrt{\frac{4mlog(1/\delta)}{(s_+ s_- m)^2}} > \epsilon^+ = \sqrt{\frac{4mlog(1/\delta)}{(\alpha s_+^2 + (1-\alpha) s_-^2)^2 m^2}} \ . \tag{16}$$

Set $\epsilon = \epsilon^-$ to have a larger step and a stricter condition, we have

$$\begin{aligned} (1+\epsilon)\mu^- &< (1-\epsilon)\mu^+ \\ \epsilon &< \frac{u^+ - u^-}{u^+ + u^-} \end{aligned} \tag{17}$$

$$\sqrt{\frac{4m \log(1/\delta)}{(s_+ s_- m)^2}} < \frac{(s_+ - s_-)^2 + (2\alpha - 1)(s_+^2 - s_-^2)}{(s_+ + s_-)^2 + (2\alpha - 1)(s_+^2 - s_-^2)}. \tag{18}$$

Because $\alpha > \frac{1}{2}$ and $s_+ > s_-$,

$$\begin{aligned} (s_+ - s_-)^2 + (2\alpha - 1)(s_+^2 - s_-^2) > 0, \\ (s_+ + s_-)^2 + (2\alpha - 1)(s_+^2 - s_-^2) > 0. \end{aligned} \tag{19}$$

So we have

$$m > \frac{4((s_+ + s_-)^2 + (2\alpha - 1)(s_+^2 - s_-^2))^2}{s_+^2 s_-^2 ((s_+ - s_-)^2 + (2\alpha - 1)(s_+^2 - s_-^2))^2}) \log(\frac{1}{\delta}). \tag{20}$$

We complete the proof by substituting δ with

$$\delta := \frac{1}{\gamma} \min((1-\alpha)s_-, \alpha(1-s_+)). \tag{21}$$

where $\gamma > 1$ is a constant.

Finally, we prove that

$$m > \frac{4((s_+ + s_-)^2 + (2\alpha - 1)(s_+^2 - s_-^2))^2}{s_+^2 s_-^2((s_+ - s_-)^2 + (2\alpha - 1)(s_+^2 - s_-^2))^2} \log\left(\frac{\gamma}{\min((1-\alpha)s_-, \alpha(1-s_+))}\right). \tag{22}$$

We can guarantee that when the above condition is met, the proposed similarity measure between two nodes from the same category is strictly larger than that for two nodes from different categories. It is able to reduce the number of noisy edges and missing edges of the single test method by $1/\gamma$ in expectation.

F Case Studies on self attention, $\mathcal{Q}$ttention and $\mathcal{B}$ttention

To better understand the effect of multiple tests (i.e. the fourth order of statistics), we illustrate the effect of self-attention, $\mathcal{Q}$**ttention** and $\mathcal{B}$**ttention** with the help of the tool BertViz[2].

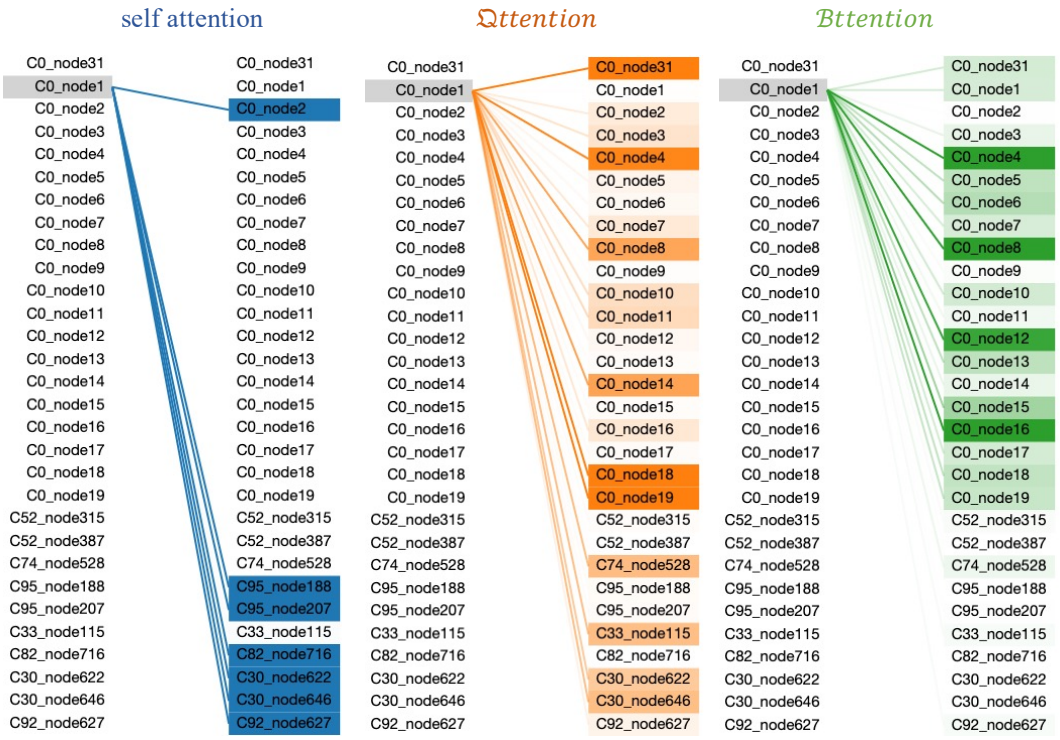


Figure 8: Case studies on self attention, $\mathcal{Q}$**ttention** and $\mathcal{B}$**ttention**. Each entry contains the category and node index, e.g. "C0_node1" means this node belongs to category 0 and the node index is 1. The shade of the colour represents the weight of attention. The darker the colour, the greater the weight. The data of this case is sampled from the last layer of $\mathcal{B}$**ttention** on the MS-Celeb part1.

As shown in the left panel of Figure 8, self-attention introduces a number of noisy connections between nodes belonging to different categories. In contrast, according to the middle panel of Figure 8, $\mathcal{Q}$**ttention** is able to introduce many more connections between nodes from the same

[2]https://github.com/jessevig/bertviz

category, and at the same time significantly reduces the weights for those noisy connections. In the last panel, the combination self-attention with the fourth order statistics, The $\mathcal{B}$**ttention** further removes those noisy connections while keeps most of the clean connections between nodes.