Contrastive General Graph Matching with Adaptive Augmentation Sampling



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Introduction

Motivation

Graph matching is crucial across various domains such as bioinformatics, social network analysis, and computer vision. The effectiveness of traditional methods, however, is curtailed by significant challenges, including those inherent in the burgeoning field of contrastive learning.



- **Dependency on Labeled Data:** Most graph matching techniques heavily rely on extensive labeled data for training. This dependence is resource-intensive and limits applicability in areas where such data is scarce or expensive to procure.
- Lack of Generalizability: Existing approaches often require additional side information or are tailored to specific graph types, which hinders their broader application. This specificity reduces the utility of graph matching methods in new or diverse areas that demand adaptable solutions.
- **Contrastive Learning Limitations:** While contrastive learning offers a

Example of visual graph matching.

promising direction for self-supervised learning in graph matching, it typically necessitates careful selection of augmentations to generate effective positive and negative samples. This requirement presents a challenge in tuning and selecting these augmentations without exacerbating the computational burden or risking model overfitting.

Experiments

Proposed Method: GCGM



Real-world Datasets						
Mathada	Pasca	al VOC	Willow	SPair-71k		
	Intsec Unfilt		Intsec	Intsec	Unfilt	
CIE (SUP)	66.8±0.4	-	82.6±0.2	69.3±0.3	_	
BBGM (SUP)	77.3 ± 0.1	$55.0 {\pm} 0.1$	96.2 ± 0.1	77.7 ± 0.2	$48.4{\pm}0.2$	
NGMv2 (SUP)	76.8 ± 0.1	56.7 ± 0.1	94.5±0.3	76.6 ± 0.2	$49.8 {\pm} 0.08$	
IPFP	45.8±0.02	31.5±0.002	80.1±0.06	57.0±0.04	31.7±0.01	
RRWM	47.2 ± 0.02	$31.7 {\pm} 0.001$	83.4±0.09	58.6 ± 0.05	$32.3 {\pm} 0.01$	
SM	46.2 ± 0.03	$30.4 {\pm} 0.002$	81.3±0.08	57.7±0.04	$30.3 {\pm} 0.01$	
GANN-GM [^]	34.5±0.3	23.4±0.2	89.3±0.1	34.7±0.4	19.4±0.3	
SCGM+BBGM	54.8 ± 0.05	<u>36.6</u> ±0.04	93.1±0.08	60.2 ± 0.05	34.1 ± 0.01	
SCGM+NGMv2	50.8 ± 0.1	32.9 ± 0.03	84.2 ± 0.1	59.8±0.1	30.5 ± 0.3	
GCGM+BBGM	56.8±0.02	36.2±0.01	<u>94.4</u> ±0.3	60.6 ± 0.1	35.9 ±0.07	
GCGM+NGMv2	57.3 ±0.11	37.4 ±0.07	95.0 ±0.1	62.6 ±0.02	35.4 ± 0.07	
Synthe	etic Dat	taset	<u> </u>	Intersect	<u>ion</u>	

Graph-centric Contrastive framework for Graph Matching (GCGM), leverages a comprehensive pool of graph augmentations to enhance robustness and effectiveness in graph matching. Boosting-inspired Adaptive Augmentation Sampler (BiAS) adaptively selects challenging augmentations tailored for graph matching, optimizing the learning process without manual tuning.

Pool of Augmentations Pairs

A distinctive feature of GCGM is its use of a large and diverse pool of graph augmentations, initially randomly initialized. This pool includes various structural and [• feature-based modifications.



Figure 2: Graph augmentation: (a) input graph; (b) node insertion; (c) node replacement; (d) edge removal. The blue node represents the inserted node, and the dotted edge indicates the added edge.

BiAS

$$v_{t+1}^i = \lambda w_t^i + (1 - \lambda) e^{\alpha (1 - \phi_t^i)}$$

Adaptive Sampling Strategy: BiAS introduces a novel weight update scheme that adaptively adjusts the probabilities of selecting specific augmentations based on their impact on the model's performance.

Weight Update Scheme: The

weights of augmentation pairs are dynamically updated based on their performance in improving the matching accuracy. Pairs that result

	Synth	netic	No outliers.		
Methods	Intsec	Unfilt	Unfiltered		
GANN-GM^	11.2 ± 0.04	10.2 ± 0.03	Includes all nodes,		
SCGM + BBGM	33.5 ± 2.0	24.3 ± 1.2	assessing robustness to		
SCGM + NGMv2	35.2 ± 0.6	25.0 ± 0.4	outliers and node count		
GCGM	58.1 ±0.5	39.9 ±0.4	variations.		

Graph Augmentation

ugmentation	Pascal VOC		Willow	SPair-71k		Synthetic	
Set	Intsec	Unfilt	Intsec	Intsec	Unfilt	Intsec	Unfilt
$\mathcal{T} \setminus \mathrm{NI}$	56.9	36.6	94.8	61.8	34.9	57.9	40.5
$\mathcal{T} \setminus \mathbf{NR}$	56.5	36.5	95.1	61.8	34.4	57.8	40.0
$\mathcal{T} \setminus \mathbf{ER}$	57.3	37.2	95.0	59.8	32.5	57.9	40.0
$\mathcal{T} \setminus \mathbf{FS}$	57.5	37.2	95.0	62.1	35.1	57.8	40.3
\mathcal{T}	57.3	37.4	95.0	62.6	35.4	58.1	39.9

Table 4: Ablation study on graph augmentations.

BiAS and Augmentation Pool

Cattings		Pascal VOC		Willow		SPair-71k			
Settings		Intsec	Unfilt	Time/h	Intsec	Time/h	Intsec	Unfilt	Time/h
Random	X	55.0	35.9	0.26	93.8	0.04	61.4	35.2	0.38
Tuning	X	55.9	36.8	23.76	94.8	4.54	61.5	35.6	31.96
Tuning + BiAS	X	55.8	37.0	23.95	95.4	4.54	61.9	36.0	31.88
Uniform	\checkmark	56.9	36.7	0.32	94.7	0.05	62.0	34.8	0.35
DIAC	1	57 2	27 4	0.20	05 0	0.05	676	25 1	0.24

Туре	Matching scenario	Hyperparameters
NI	node outlier (unequal node count in two views)	$p_{ni} \in [0.1, 0.9]$: fraction of nodes inserted; $k_{ni} \ge 2$: size of subset; $aggr_{ni} \in \{\text{mean, max}\}$: aggregation function; $e_{ni} \ge 1$: number of edges inserted
NR	node outlier (equal node count in two views)	$p_{nr} \in [0.1, 0.9]$: fraction of nodes replaced; $k_{nr} \ge 2$: size of subset; $aggr_{nr} \in \{\text{mean, max}\}$: aggregation function; $e_{nr} \ge 1$: number of edges inserted
ER	sparse/noisy first- order connections	$p_{\rm er} \in [0.1, 0.9]$: probability of each edge being removed
FS	feature variations & noises	$\alpha \in [0.2, 0.8]: \text{ lower bound of uniform distribution;} \\ \beta \in [1.2, 1.8]: \text{ upper bound of uniform distribution}$

Table 1: Details of the four major types of graph augmentation.

in lower performance scores are frequently sampled more in subsequent training iterations, encouraging the model to focus on more challenging and informative examples.

 $\lambda = 0.6, \alpha = 2$

w _t	ϕ_t	w_{t+1}
1	0.9	1.09
1	0.4	1.93

Example of weight update.

DIAS V 51.5 51.4 0.39 | 93.0 | 0.03 | 02.0 | 33.4 | 0.34

Table 5: Performance of different initialization of augmentations and the use of augmentation pool. Time (hour) represents the total wall clock time spent on tuning the augmentations (for 'Tuning' methods) and training the model. The ' \mathcal{P} ' column indicates if a pool of augmentation pairs is used.

