

DShield: Defending against Backdoor Attacks on Graph Neural Networks via Discrepancy Learning

Hao Yu¹, Chuan Ma², Xinhang Wan¹, Jun Wang¹, Tao Xiang², Meng Shen³, and Xinwang Liu¹

National University of Defense Technology

Chongqing University

Beijing Institute of Technology

Speaker: **Zhongming Wang**



国防科技大学
NATIONAL UNIVERSITY
OF DEFENSE TECHNOLOGY



重慶大學
CHONGQING UNIVERSITY



北京理工大學
BEIJING INSTITUTE OF TECHNOLOGY

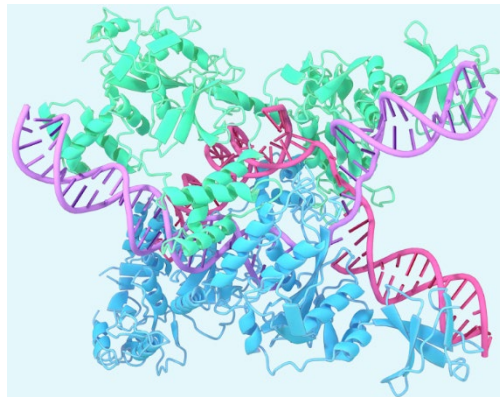
Feb. 27, 2025@San Diego, USA

Graph Neural Networks

- Many types of data can be represented as **graph structures**, e.g., social networks and protein molecules
- GNNs **propagate** information across graph structures, effectively capturing complex relationships between nodes and edges
- Tasks: node classification, graph classification, link prediction



Social Networks



Drug Discovery



Smart City



Bank Risk Management

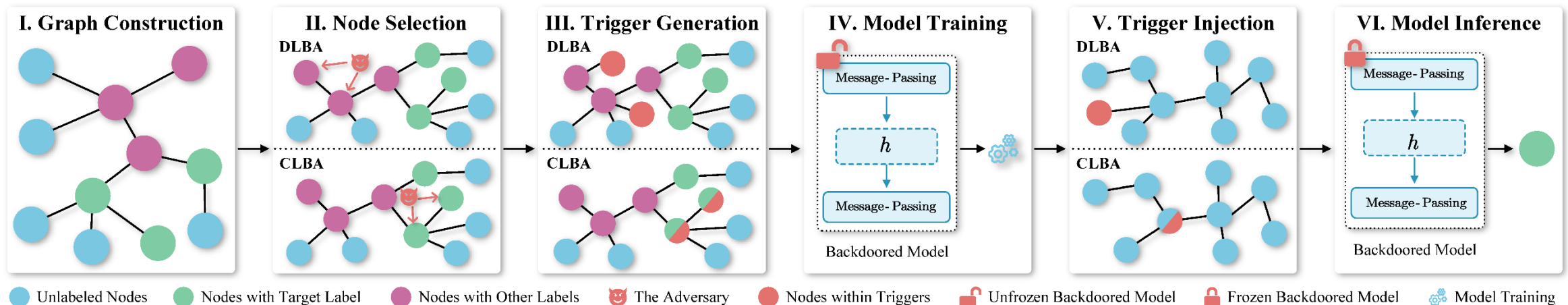
Backdoor Attacks on GNNs

➤ Backdoor attacks

- Inducing **undesirable** behaviors in **backdoored** models by injecting **triggers** into original graphs

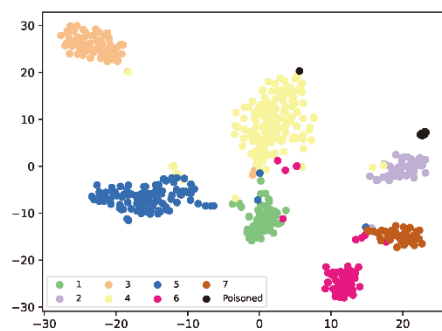
➤ Two variants

- Dirty-label backdoor attacks (**DLBAs**): Altering label information
- Clean-label backdoor attacks (**CLBAs**): Altering attributes of normal nodes

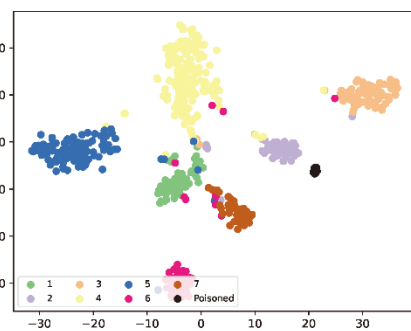


➤ Two phenomena

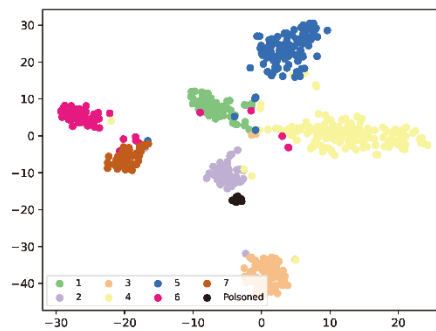
- Semantic Drift of DLBAs: between node's attributes and structures, and their labels
- Attribute over-emphasis of CLBAs: high similarities of important attributes



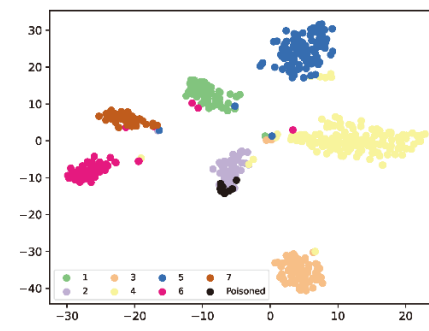
(a)



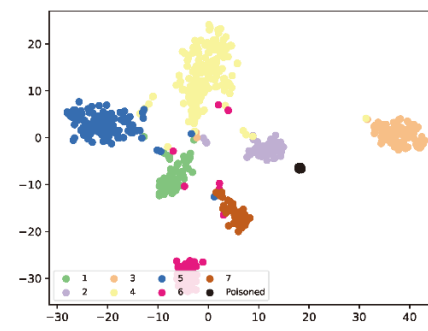
(b)



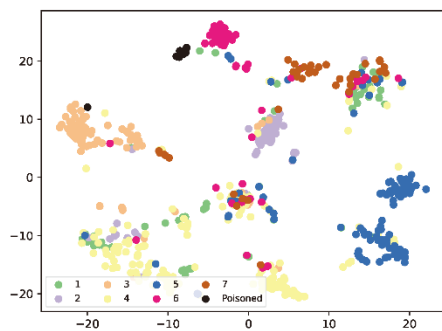
(c)



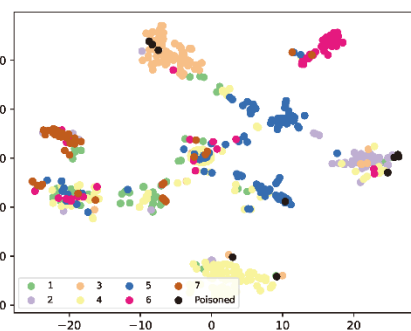
(a)



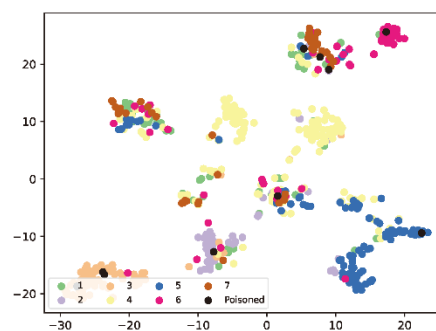
(b)



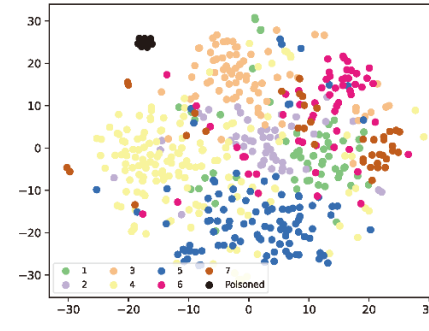
(d)



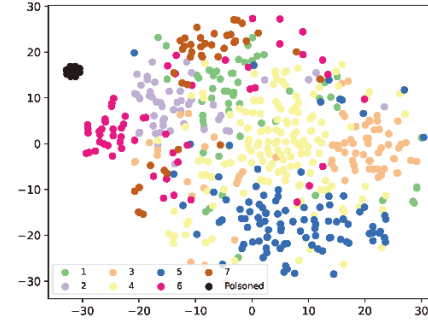
(e)



(f)



(c)



(d)

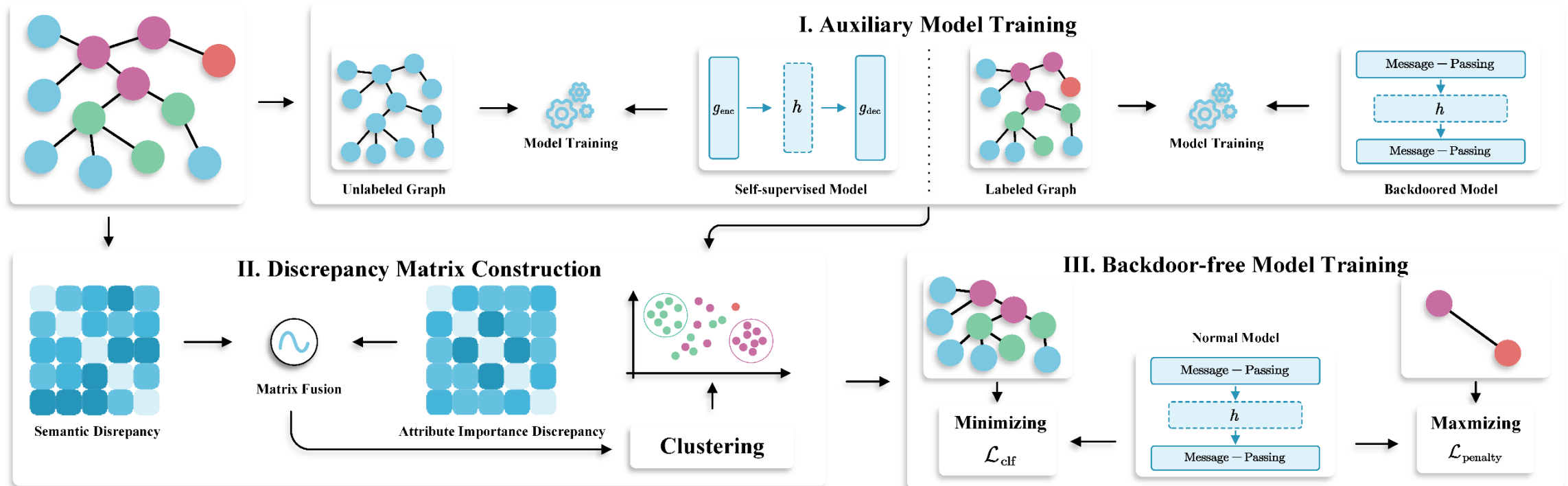
Semantic Drift

Attribute Over-emphasis

Proposed DShield

➤ Pipeline

- Auxiliary Model Training: Preparing to identify poisoned nodes in DLBAs
- Discrepancy Matrix Construction: Detecting poisoned nodes in both DLBAs and CLBAs
- Backdoor-free Model Training: Training a model without backdoors

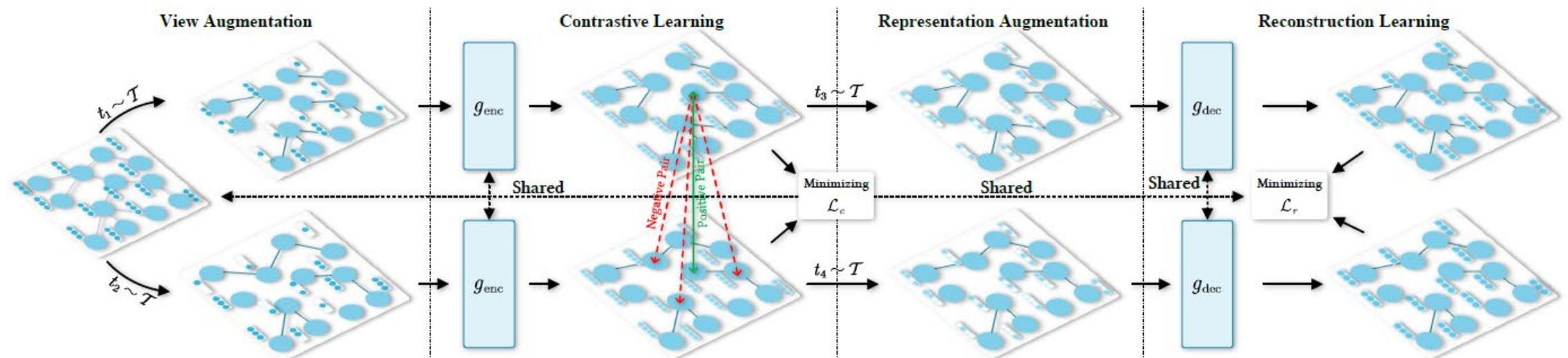


➤ Backdoored Model

$$\mathcal{L} = \frac{1}{|\mathcal{V}^{train}|} \sum_{v_i \in \mathcal{V}^{train}} \text{CE}(\tilde{f}(\mathbf{A}, \mathbf{X})_i, \mathbf{y}_i)$$

➤ Self-supervised Model

- View Augmentation: Sampling two stochastic augmentation functions
- View Encoding: Extracting latent representations of nodes
- Contrast and Reconstruction: Distinguishing between representations of two nodes



➤ View Augmentation

$$P\{(v_i, v_j) \in \hat{\mathcal{E}}\} = 1 - p_{ij}^s; \quad p_{ij}^s = \frac{\exp(-s(\tilde{\mathbf{h}}_i, \tilde{\mathbf{h}}_j))}{\sum_{(v_k, v_t) \in \tilde{\mathcal{E}}} \exp(-s(\tilde{\mathbf{h}}_k, \tilde{\mathbf{h}}_t))}$$

Structure-level

$$\mathbf{g}_i = \frac{\partial \text{CE}(f(\tilde{\mathbf{A}}, \tilde{\mathbf{X}}; \phi)_i, \mathbf{y}_i)}{\partial \mathbf{x}_i}; \quad \hat{\mathbf{A}} = [\mathbf{x}_1 \odot \hat{\mathbf{m}}_1; \dots; \mathbf{x}_N \odot \hat{\mathbf{m}}_N]^\top$$

Attribute-level

➤ View Encoding

$$\hat{\mathbf{H}}_1 = g_{\text{enc}}(\hat{\mathbf{A}}_1, \hat{\mathbf{X}}_1), \quad \hat{\mathbf{H}}_2 = g_{\text{enc}}(\hat{\mathbf{A}}_2, \hat{\mathbf{X}}_2)$$

➤ Contrast and Reconstruction

Contrast

$$\mathcal{L}_{c_1} = \sum_{v_i \in \mathcal{V}^{\text{train}}} - \frac{1}{|\mathcal{P}(v_i)|} \sum_{v_j \in \mathcal{P}(v_i)} \log \frac{\exp(s(\hat{\mathbf{h}}_{1,i}, \hat{\mathbf{h}}_{2,j})/\tau)}{\sum_{v_k \in \mathcal{N}(v_i)} \exp(s(\hat{\mathbf{h}}_{1,i}, \hat{\mathbf{h}}_{2,k})/\tau)};$$

Reconstruction

$$\hat{\mathbf{H}}'_1 = [\hat{\mathbf{h}}_1 \odot \hat{\mathbf{m}}'_1; \hat{\mathbf{h}}_2 \odot \hat{\mathbf{m}}'_2; \dots; \hat{\mathbf{h}}_N \odot \hat{\mathbf{m}}'_N]^\top \quad \hat{\mathbf{H}}'_1 = g_{\text{dec}}(\hat{\mathbf{A}}_1, \hat{\mathbf{X}}'_1), \quad \hat{\mathbf{H}}'_2 = g_{\text{dec}}(\hat{\mathbf{A}}_2, \hat{\mathbf{X}}'_2)$$

$$\mathcal{L}_r = \frac{1}{2} \sum_{i=1}^N (1 - s(\tilde{\mathbf{x}}_i, \hat{\mathbf{x}}'_{1,i}))^2 + \frac{1}{2} \sum_{i=1}^N (1 - s(\tilde{\mathbf{x}}_i, \hat{\mathbf{x}}'_{2,i}))^2$$

Discrepancy Matrix Construction

➤ Semantic Discrepancy Matrix

$$\widetilde{\mathbf{H}}_{\text{sl}} = \widetilde{f}(\widetilde{\mathbf{A}}, \widetilde{\mathbf{X}}; \phi_1), \quad \widetilde{\mathbf{H}}_{\text{ssl}} = g_{\text{enc}}(\widetilde{\mathbf{A}}, \widetilde{\mathbf{X}})$$

$$\mathbf{D}_{\text{sl}}^y = \mathbf{1} \text{diag}(\widetilde{\mathbf{H}}_{\text{sl}}^y \widetilde{\mathbf{H}}_{\text{sl}}^{y\top})^\top - 2\widetilde{\mathbf{H}}_{\text{sl}}^y \widetilde{\mathbf{H}}_{\text{sl}}^{y\top} + \text{diag}(\widetilde{\mathbf{H}}_{\text{sl}}^y \widetilde{\mathbf{H}}_{\text{sl}}^{y\top}) \mathbf{1}^\top;$$

$$\mathbf{D}_{\text{ssl}}^y = \mathbf{1} \text{diag}(\widetilde{\mathbf{H}}_{\text{ssl}}^y \widetilde{\mathbf{H}}_{\text{ssl}}^{y\top})^\top - 2\widetilde{\mathbf{H}}_{\text{ssl}}^y \widetilde{\mathbf{H}}_{\text{ssl}}^{y\top} + \text{diag}(\widetilde{\mathbf{H}}_{\text{ssl}}^y \widetilde{\mathbf{H}}_{\text{ssl}}^{y\top}) \mathbf{1}^\top,$$

$$\mathbf{D}_s^y = \max(\mathbf{D}_{\text{ssl}}^y - \mathbf{D}_{\text{sl}}^y, 0)$$

➤ Attribute Importance Discrepancy Matrix

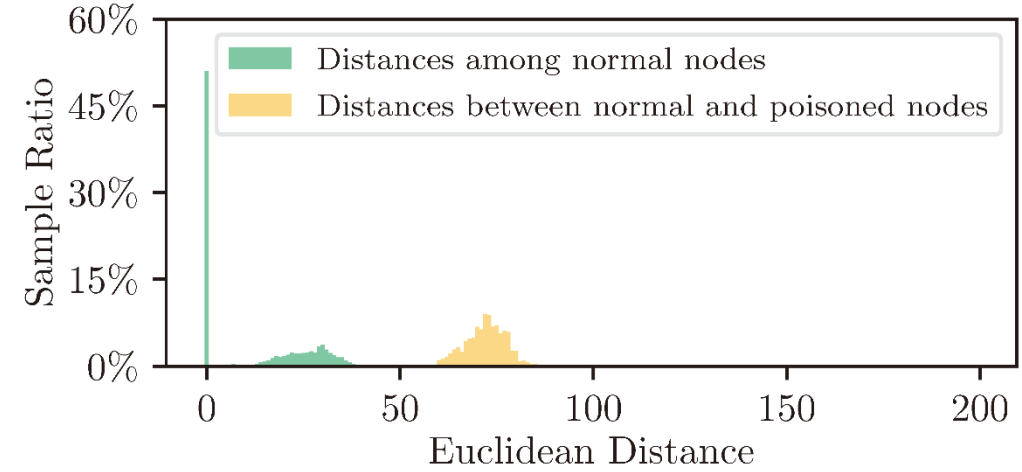
$$\mathbf{g}_i = \frac{\partial \text{CE}(f(\widetilde{\mathbf{A}}, \widetilde{\mathbf{X}}; \phi)_i, \mathbf{y}_i)}{\partial \mathbf{x}_i}, \quad w_{ij} = \mathbb{1}_{g_{ij} > 0} \quad \widetilde{\mathbf{X}}^y = \widetilde{\mathbf{X}}^y \circ \mathbf{W}^y$$

$$\overline{\mathbf{X}}^y = \text{UMAP}(\widetline{\mathbf{X}}^y, d)$$

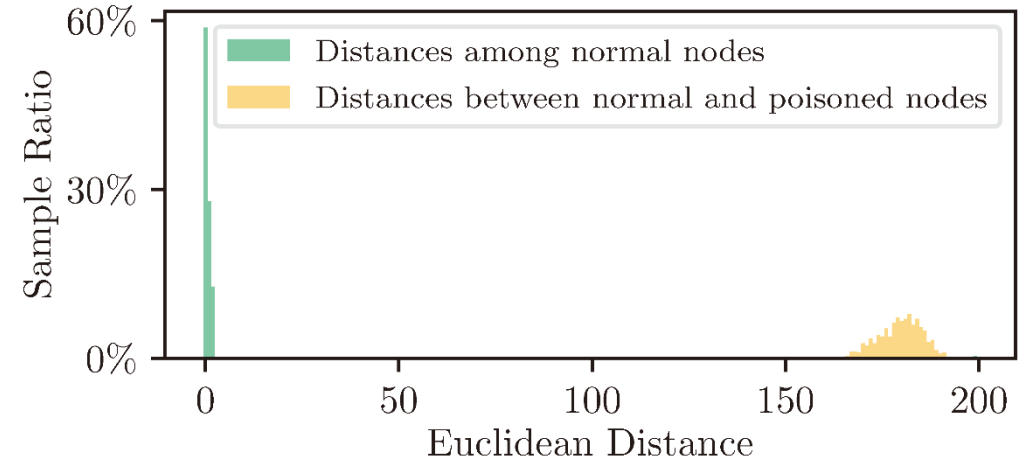
$$\mathbf{D}_a^y = \mathbf{1} \text{diag}(\overline{\mathbf{X}}^y \overline{\mathbf{X}}^{y\top})^\top - 2\overline{\mathbf{X}}^y \overline{\mathbf{X}}^{y\top} + \text{diag}(\overline{\mathbf{X}}^y \overline{\mathbf{X}}^{y\top}) \mathbf{1}^\top$$

➤ Clustering

$$\mathbf{D}_y = \frac{\text{std}(\mathbf{D}_s^y)}{\text{std}(\mathbf{D}_s^y) + \text{std}(\mathbf{D}_a^y)} \mathbf{D}_s^y + \frac{\text{std}(\mathbf{D}_a^y)}{\text{std}(\mathbf{D}_s^y) + \text{std}(\mathbf{D}_a^y)} \mathbf{D}_a^y; \quad \mathcal{V}_1, \mathcal{V}_2 = \text{HDBSCAN}(\mathbf{D}_y)$$



(a) Before Dimensionality Reduction



(b) After Dimensionality Reduction

Importance of UMAP

➤ Target Label Discovery

● Model Training

$$\tilde{A}_{ij}^{y, \text{pos}} \begin{cases} = \tilde{A}_{ij}, & \text{if } v_i \notin \mathcal{V}_2^y \text{ and } v_j \notin \mathcal{V}_2^y \\ = 0, & \text{if } v_i \in \mathcal{V}_2^y \text{ or } v_j \in \mathcal{V}_2^y \end{cases}$$

$$\tilde{A}_{ij}^{y, \text{neg}} \begin{cases} = \tilde{A}_{ij}, & \text{if } v_i \in \mathcal{V}_2^y \text{ or } v_j \in \mathcal{V}_2^y \\ = 0, & \text{if } v_i \notin \mathcal{V}_2^y \text{ and } v_j \notin \mathcal{V}_2^y \end{cases}$$

$$\mathcal{L}_{\text{pos}} = \frac{1}{|\mathcal{V}_1^y|} \sum_{v_i \in \mathcal{V}_1^y} \text{CE}(f'(\tilde{\mathbf{A}}^{y, \text{pos}}, \tilde{\mathbf{X}})_i, \mathbf{y}_i) \quad \mathcal{L}_{\text{neg}} = \frac{1}{|\mathcal{V}_2^y|} \sum_{v_i \in \mathcal{V}_2^y} \text{CE}(f'(\tilde{\mathbf{A}}^{y, \text{neg}}, \tilde{\mathbf{X}})_i, \mathbf{y}_i). \quad \mathcal{L} = \mathcal{L}_{\text{pos}} - \log(\mathcal{L}_{\text{neg}})$$

● Target Label

$$\mathbf{h}^y = \frac{1}{|\mathcal{V}_1^y|} \sum_{v_i \in \mathcal{V}_1^y} f'(\tilde{\mathbf{A}}^{y, \text{pos}}, \tilde{\mathbf{X}}; \phi'_1)_i \quad \mathbf{s}^y = \frac{1}{|\mathcal{V}_2^y|} \sum_{v_i \in \mathcal{V}_2^y} \|f'(\tilde{\mathbf{A}}^{y, \text{neg}}, \tilde{\mathbf{X}}; \phi'_1)_i - \mathbf{h}^y\|_2^2$$

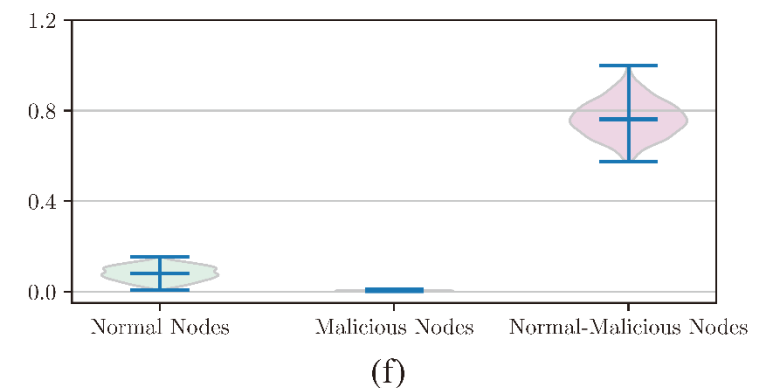
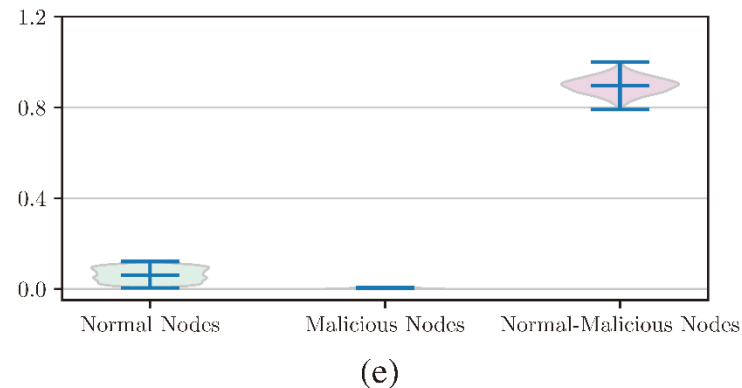
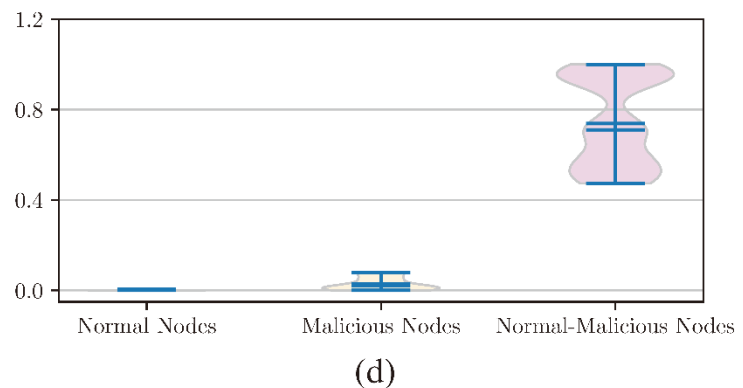
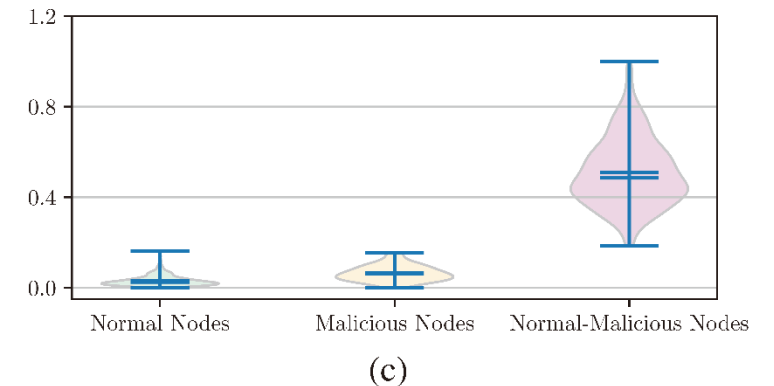
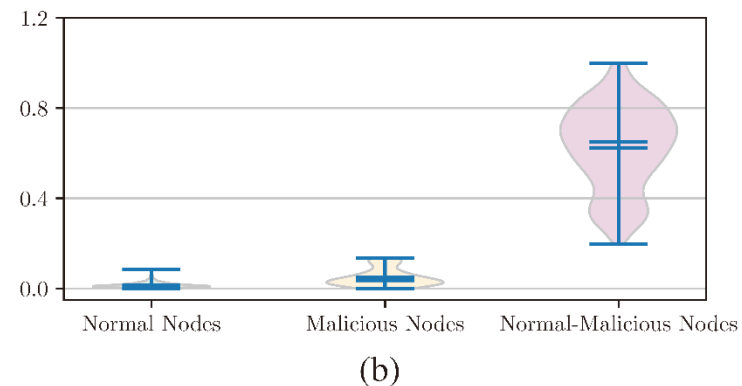
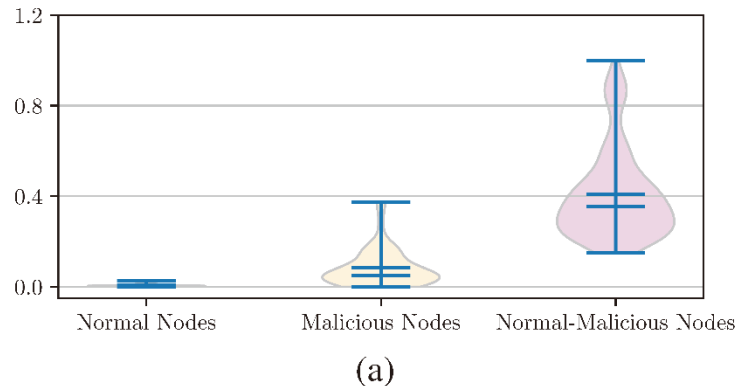
$$\mathcal{Y} = \left\{ y \mid y \in [1, C] \wedge \frac{|s^y - \text{median}(\mathbf{s})|}{\text{mad}(\mathbf{s})} \leq \beta \right\}$$

➤ Backdoor-free Training

$$\mathcal{L}_{\text{clf}} = \frac{1}{|\mathcal{V}^\dagger|} \sum_{v_i \in \mathcal{V}^\dagger} \text{CE}(f(\tilde{\mathbf{A}}, \tilde{\mathbf{X}})_i, \mathbf{y}_i) \quad \mathcal{L}_{\text{penalty}} = \frac{1}{|\mathcal{V}^\dagger|} \sum_{v_i \in \mathcal{V}^\dagger} \text{CE}(f(\tilde{\mathbf{A}}', \tilde{\mathbf{X}})_i, \mathbf{y}_i) \quad \mathcal{L} = \mathcal{L}_{\text{clf}} - \gamma \log \mathcal{L}_{\text{penalty}}$$

➤ Elements of Discrepancy Matrices

- Differences in elements of two discrepancy matrices between normal and malicious nodes in the Cora dataset.



➤ Seven DLBAs & Four Datasets

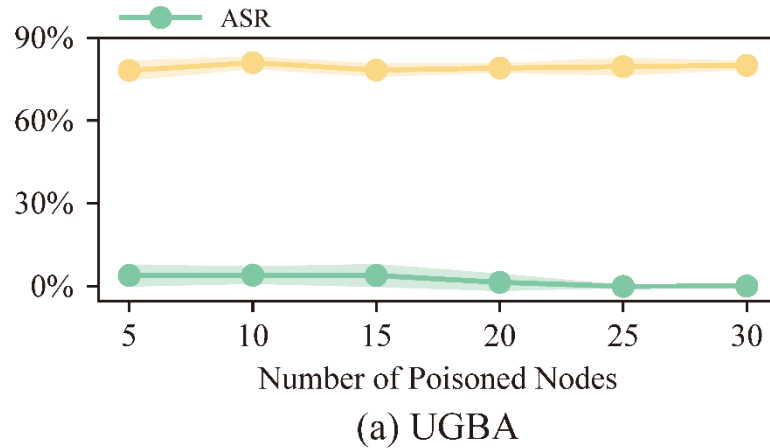
Datasets	Defenses	SBA [50]		GTA [38]		EBA [43]		GB-FGSM [4]		LGCB [4]		UGBA [5]		TRAP [44]	
		ASR ↓	ACC ↑	ASR ↓	ACC ↑	ASR ↓	ACC ↑	ASR ↓	ACC ↑	ASR ↓	ACC ↑	ASR ↓	ACC ↑	ASR ↓	ACC ↑
Cora	<i>no-defense</i>	53.14 $\pm$ 2.10	84.07 $\pm$ 1.02	96.22 $\pm$ 4.41	82.74 $\pm$ 2.38	71.81 $\pm$ 9.44	83.48 $\pm$ 1.32	97.34 $\pm$ 1.64	82.52 $\pm$ 1.53	97.79 $\pm$ 1.01	83.70 $\pm$ 0.52	97.51 $\pm$ 2.07	83.03 $\pm$ 1.32	19.86 $\pm$ 2.42	84.15 $\pm$ 1.09
	Prune [5]	30.26 $\pm$ 2.78	83.26 $\pm$ 0.96	16.42 $\pm$ 2.35	84.00 $\pm$ 1.80	100.0 $\pm$ 0.00	80.74 $\pm$ 1.57	56.98 $\pm$ 1.00	81.92 $\pm$ 3.36	56.98 $\pm$ 0.93	81.92 $\pm$ 1.95	54.47 $\pm$ 2.19	83.41 $\pm$ 1.52	11.36 $\pm$ 2.08	81.48 $\pm$ 1.87
	Isolate [5]	04.06 $\pm$ 2.42	37.19 $\pm$ 1.07	03.54 $\pm$ 1.85	77.78 $\pm$ 1.48	96.43 $\pm$ 1.66	80.67 $\pm$ 1.29	56.09 $\pm$ 0.78	81.85 $\pm$ 1.08	56.46 $\pm$ 0.78	81.63 $\pm$ 1.97	56.02 $\pm$ 0.80	80.74 $\pm$ 3.21	10.15 $\pm$ 1.94	67.26 $\pm$ 1.63
	PXGBD [9]	46.37 $\pm$ 2.03	84.00 $\pm$ 0.85	97.05 $\pm$ 2.46	83.40 $\pm$ 1.74	71.04 $\pm$ 9.65	82.74 $\pm$ 1.40	97.86 $\pm$ 1.44	82.74 $\pm$ 0.93	98.38 $\pm$ 1.64	82.37 $\pm$ 1.13	97.86 $\pm$ 2.17	84.00 $\pm$ 0.55	22.14 $\pm$ 2.31	81.55 $\pm$ 1.37
	GXGBD [9]	56.28 $\pm$ 1.31	84.17 $\pm$ 1.30	83.77 $\pm$ 1.56	83.42 $\pm$ 1.10	88.19 $\pm$ 2.81	73.52 $\pm$ 2.75	98.16 $\pm$ 1.16	84.07 $\pm$ 1.09	98.43 $\pm$ 1.01	84.26 $\pm$ 0.98	80.81 $\pm$ 2.30	83.15 $\pm$ 2.31	19.70 $\pm$ 3.84	81.18 $\pm$ 0.80
	ABL [21]	54.86 $\pm$ 2.37	82.81 $\pm$ 1.40	97.79 $\pm$ 2.09	82.59 $\pm$ 2.38	81.43 $\pm$ 3.98	83.26 $\pm$ 1.37	97.56 $\pm$ 1.34	83.11 $\pm$ 0.81	98.08 $\pm$ 0.84	83.70 $\pm$ 0.59	97.42 $\pm$ 1.04	83.89 $\pm$ 2.19	18.38 $\pm$ 2.52	83.33 $\pm$ 0.64
	RS [49]	08.12 $\pm$ 1.51	80.81 $\pm$ 1.03	94.83 $\pm$ 2.93	80.67 $\pm$ 0.99	58.92 $\pm$ 2.72	81.11 $\pm$ 1.05	97.27 $\pm$ 0.85	80.15 $\pm$ 1.24	96.97 $\pm$ 1.09	80.15 $\pm$ 1.84	90.41 $\pm$ 1.56	80.81 $\pm$ 1.03	20.81 $\pm$ 1.78	79.85 $\pm$ 1.13
	DShield	01.33 $\pm$ 2.56	81.78 $\pm$ 1.93	00.74 $\pm$ 0.74	81.92 $\pm$ 1.98	02.95 $\pm$ 2.32	81.50 $\pm$ 1.08	02.51 $\pm$ 2.83	82.29 $\pm$ 0.71	00.89 $\pm$ 1.15	82.57 $\pm$ 0.51	01.33 $\pm$ 1.21	82.15 $\pm$ 0.80	13.38 $\pm$ 2.12	82.07 $\pm$ 1.65
PubMed	<i>no-defense</i>	62.84 $\pm$ 5.05	85.17 $\pm$ 0.20	96.54 $\pm$ 2.28	85.09 $\pm$ 0.26	84.79 $\pm$ 2.10	85.23 $\pm$ 0.17	99.67 $\pm$ 0.66	85.03 $\pm$ 0.15	97.14 $\pm$ 1.44	85.16 $\pm$ 0.22	93.27 $\pm$ 3.23	85.20 $\pm$ 0.25	48.34 $\pm$ 0.93	85.21 $\pm$ 0.38
	Prune [5]	53.01 $\pm$ 2.59	85.33 $\pm$ 0.37	46.15 $\pm$ 0.40	85.30 $\pm$ 0.21	99.95 $\pm$ 0.00	85.44 $\pm$ 0.18	70.28 $\pm$ 0.40	85.31 $\pm$ 0.25	69.36 $\pm$ 0.56	85.36 $\pm$ 0.18	66.85 $\pm$ 1.47	85.38 $\pm$ 0.29	41.77 $\pm$ 0.28	85.33 $\pm$ 0.25
	Isolate [5]	40.69 $\pm$ 0.47	81.25 $\pm$ 0.28	45.46 $\pm$ 0.22	84.47 $\pm$ 0.21	99.36 $\pm$ 0.38	84.56 $\pm$ 0.22	70.16 $\pm$ 0.31	85.14 $\pm$ 0.15	69.80 $\pm$ 0.90	85.11 $\pm$ 0.13	67.62 $\pm$ 1.34	85.14 $\pm$ 0.20	43.30 $\pm$ 3.41	83.34 $\pm$ 1.59
	PXGBD [9]	55.26 $\pm$ 3.06	83.55 $\pm$ 1.01	99.67 $\pm$ 0.54	83.17 $\pm$ 0.92	86.26 $\pm$ 5.89	82.64 $\pm$ 1.57	100.0 $\pm$ 0.00	82.32 $\pm$ 1.47	100.0 $\pm$ 0.00	82.38 $\pm$ 0.99	91.80 $\pm$ 0.87	82.61 $\pm$ 1.35	49.80 $\pm$ 1.70	81.98 $\pm$ 1.45
	GXGBD [9]	56.20 $\pm$ 1.18	83.93 $\pm$ 0.76	99.71 $\pm$ 0.35	82.57 $\pm$ 0.64	90.21 $\pm$ 2.27	75.45 $\pm$ 2.23	100.0 $\pm$ 0.00	82.31 $\pm$ 0.85	100.0 $\pm$ 0.00	82.41 $\pm$ 0.82	81.95 $\pm$ 0.00	82.82 $\pm$ 0.91	49.36 $\pm$ 3.16	82.84 $\pm$ 1.47
	ABL [21]	65.16 $\pm$ 4.55	85.21 $\pm$ 0.38	99.98 $\pm$ 0.03	85.13 $\pm$ 0.21	89.08 $\pm$ 2.61	84.98 $\pm$ 0.16	99.98 $\pm$ 0.03	84.89 $\pm$ 0.13	100.0 $\pm$ 0.00	85.04 $\pm$ 0.17	91.45 $\pm$ 5.44	85.00 $\pm$ 0.30	49.36 $\pm$ 0.68	84.64 $\pm$ 1.01
	RS [49]	51.10 $\pm$ 3.63	83.13 $\pm$ 0.45	97.53 $\pm$ 1.33	82.99 $\pm$ 0.37	89.86 $\pm$ 1.20	83.68 $\pm$ 0.32	100.0 $\pm$ 0.00	83.39 $\pm$ 0.58	100.0 $\pm$ 0.00	83.40 $\pm$ 0.57	59.68 $\pm$ 4.13	83.17 $\pm$ 0.31	48.48 $\pm$ 3.50	82.83 $\pm$ 0.71
	DShield	35.91 $\pm$ 3.43	82.01 $\pm$ 0.47	00.73 $\pm$ 0.33	85.28 $\pm$ 0.12	02.78 $\pm$ 1.58	85.02 $\pm$ 0.29	00.66 $\pm$ 0.72	83.71 $\pm$ 0.24	03.54 $\pm$ 1.97	84.11 $\pm$ 0.56	02.24 $\pm$ 0.97	84.70 $\pm$ 0.42	45.77 $\pm$ 0.76	82.17 $\pm$ 0.36

Two CLBAs & Four Datasets

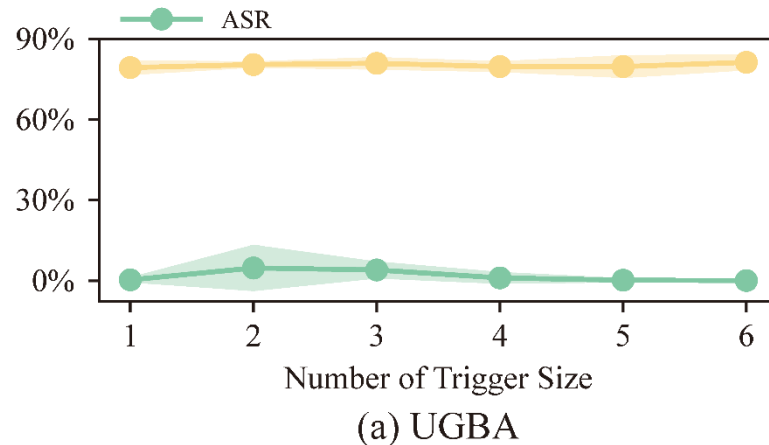
Defenses	Cora				PubMed				Flickr				OGBN-arXiv			
	GCBA [49]		PerCBA [45]		GCBA [49]		PerCBA [45]		GCBA [49]		PerCBA [45]		GCBA [49]		PerCBA [45]	
	ASR ↓	ACC ↑	ASR ↓	ACC ↑	ASR ↓	ACC ↑	ASR ↓	ACC ↑	ASR ↓	ACC ↑	ASR ↓	ACC ↑	ASR ↓	ACC ↑	ASR ↓	ACC ↑
<i>no-defense</i>	81.74 $\pm$ 3.93	84.74 $\pm$ 1.27	84.87 $\pm$ 3.13	83.18 $\pm$ 1.54	78.86 $\pm$ 7.76	85.30 $\pm$ 0.13	100.0 $\pm$ 0.00	85.23 $\pm$ 0.13	60.48 $\pm$ 4.72	50.84 $\pm$ 0.16	07.05 $\pm$ 5.11	50.86 $\pm$ 0.18	26.84 $\pm$ 2.79	60.43 $\pm$ 0.30	18.06 $\pm$ 1.48	59.81 $\pm$ 0.15
Prune [5]	99.70 $\pm$ 0.31	82.37 $\pm$ 0.42	69.01 $\pm$ 1.31	82.66 $\pm$ 0.55	99.83 $\pm$ 0.18	85.55 $\pm$ 0.16	95.58 $\pm$ 4.50	85.34 $\pm$ 0.30	78.50 $\pm$ 3.63	49.97 $\pm$ 0.27	30.38 $\pm$ 2.70	50.12 $\pm$ 0.15	23.14 $\pm$ 0.30	60.40 $\pm$ 0.35	19.03 $\pm$ 1.15	59.78 $\pm$ 0.17
Isolate [5]	99.88 $\pm$ 0.21	71.92 $\pm$ 2.05	81.73 $\pm$ 2.35	69.04 $\pm$ 2.07	99.10 $\pm$ 1.09	84.12 $\pm$ 0.14	97.88 $\pm$ 1.05	84.26 $\pm$ 1.01	66.44 $\pm$ 5.40	50.34 $\pm$ 0.08	30.15 $\pm$ 3.17	50.52 $\pm$ 0.18	22.24 $\pm$ 0.98	60.43 $\pm$ 0.08	19.00 $\pm$ 0.64	59.75 $\pm$ 0.18
PXGBD [8]	82.29 $\pm$ 3.66	82.89 $\pm$ 0.88	27.68 $\pm$ 2.58	82.74 $\pm$ 1.19	96.91 $\pm$ 3.08	82.82 $\pm$ 1.34	93.11 $\pm$ 0.36	83.31 $\pm$ 1.54	26.50 $\pm$ 0.79	47.74 $\pm$ 0.98	00.16 $\pm$ 0.28	47.29 $\pm$ 2.67	11.87 $\pm$ 0.71	60.42 $\pm$ 0.17	00.78 $\pm$ 0.49	60.42 $\pm$ 0.89
GXGBD [8]	04.61 $\pm$ 1.77	82.96 $\pm$ 0.52	80.69 $\pm$ 2.98	84.30 $\pm$ 1.07	97.30 $\pm$ 0.51	82.43 $\pm$ 1.11	72.90 $\pm$ 1.18	82.78 $\pm$ 0.32	49.90 $\pm$ 2.39	40.57 $\pm$ 0.30	02.42 $\pm$ 4.19	40.61 $\pm$ 0.44	09.21 $\pm$ 1.54	55.94 $\pm$ 0.16	00.34 $\pm$ 0.14	48.51 $\pm$ 0.43
ABL [20]	79.70 $\pm$ 3.89	84.30 $\pm$ 1.35	88.68 $\pm$ 4.25	83.70 $\pm$ 0.91	83.40 $\pm$ 1.46	84.98 $\pm$ 0.23	82.50 $\pm$ 3.53	85.13 $\pm$ 0.36	66.88 $\pm$ 0.84	50.49 $\pm$ 0.21	01.35 $\pm$ 0.60	50.68 $\pm$ 0.04	02.73 $\pm$ 3.84	58.40 $\pm$ 1.61	22.04 $\pm$ 2.39	60.23 $\pm$ 0.79
RS [49]	42.44 $\pm$ 4.11	81.26 $\pm$ 1.92	85.22 $\pm$ 3.15	80.74 $\pm$ 1.14	74.05 $\pm$ 1.70	83.28 $\pm$ 0.58	88.46 $\pm$ 3.41	83.51 $\pm$ 0.27	00.02 $\pm$ 0.03	40.48 $\pm$ 0.09	23.72 $\pm$ 0.59	40.50 $\pm$ 0.16	27.76 $\pm$ 0.08	55.36 $\pm$ 0.10	43.48 $\pm$ 5.32	55.38 $\pm$ 0.19
DSshield	01.11 $\pm$ 1.04	81.18 $\pm$ 1.32	03.41 $\pm$ 2.53	81.63 $\pm$ 2.04	01.91 $\pm$ 1.66	84.24 $\pm$ 0.24	08.09 $\pm$ 2.04	83.69 $\pm$ 2.01	01.66 $\pm$ 2.34	50.42 $\pm$ 0.25	00.04 $\pm$ 0.05	50.86 $\pm$ 0.06	00.00 $\pm$ 0.00	59.96 $\pm$ 0.54	00.00 $\pm$ 0.00	59.92 $\pm$ 0.44

- DSshield exhibits a **notable** performance against most backdoor attacks, surpassing the efficacy of conventional defenses
- Although DSshield demonstrates superior defense performance in most scenarios, a **slight decline** in model performance on normal nodes is observed

➤ Six Poisoning Rates



➤ Six Trigger Sizes

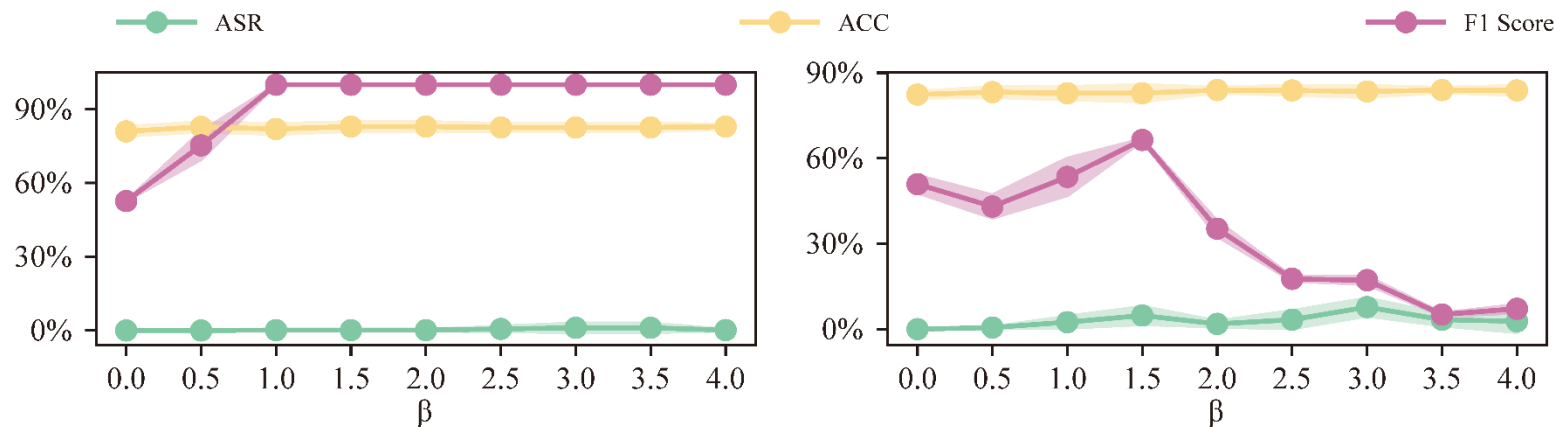


➤ Five Adaptive Attacks

Datasets	Attacks	<i>no-defense</i>		<i>no-poison</i>		DShield	
		ASR ↓	ACC ↑	ASR ↓	ACC ↑	ASR ↓	ACC ↑
Cora	UGBA+LGCB	97.91 $\pm$ 0.93	84.32 $\pm$ 1.19	08.00 $\pm$ 0.21	78.02 $\pm$ 5.00	00.49 $\pm$ 4.16	82.22 $\pm$ 6.72
	UGBA+GCBA	88.07 $\pm$ 1.89	83.83 $\pm$ 2.04	08.00 $\pm$ 0.43	78.15 $\pm$ 4.98	03.94 $\pm$ 4.49	81.97 $\pm$ 5.46
	GCBA+PerCBA	74.78 $\pm$ 3.77	83.82 $\pm$ 1.13	08.24 $\pm$ 0.93	77.53 $\pm$ 3.36	08.49 $\pm$ 2.82	80.99 $\pm$ 4.64
	AdaDA	84.51 $\pm$ 2.61	84.07 $\pm$ 0.52	07.75 $\pm$ 0.64	84.07 $\pm$ 1.29	04.43 $\pm$ 3.13	76.86 $\pm$ 0.26
	AdaCA	89.85 $\pm$ 6.52	82.59 $\pm$ 0.52	09.47 $\pm$ 5.18	84.69 $\pm$ 0.94	00.74 $\pm$ 0.98	81.23 $\pm$ 2.46
PubMed	UGBA+LGCB	88.73 $\pm$ 5.16	84.98 $\pm$ 0.26	37.92 $\pm$ 1.71	82.23 $\pm$ 1.97	00.15 $\pm$ 0.27	82.50 $\pm$ 0.74
	UGBA+GCBA	91.89 $\pm$ 3.07	85.42 $\pm$ 0.03	39.61 $\pm$ 4.17	83.66 $\pm$ 0.43	02.92 $\pm$ 1.17	82.51 $\pm$ 1.19
	GCBA+PerCBA	89.01 $\pm$ 5.25	84.53 $\pm$ 0.62	39.52 $\pm$ 3.38	83.49 $\pm$ 0.61	13.39 $\pm$ 2.08	81.28 $\pm$ 1.17
	AdaDA	88.72 $\pm$ 3.34	85.22 $\pm$ 0.04	40.23 $\pm$ 0.72	83.17 $\pm$ 3.57	32.61 $\pm$ 8.89	79.17 $\pm$ 1.47
	AdaCA	87.88 $\pm$ 5.12	85.13 $\pm$ 0.31	03.82 $\pm$ 1.68	85.39 $\pm$ 0.13	05.68 $\pm$ 0.79	82.60 $\pm$ 1.87

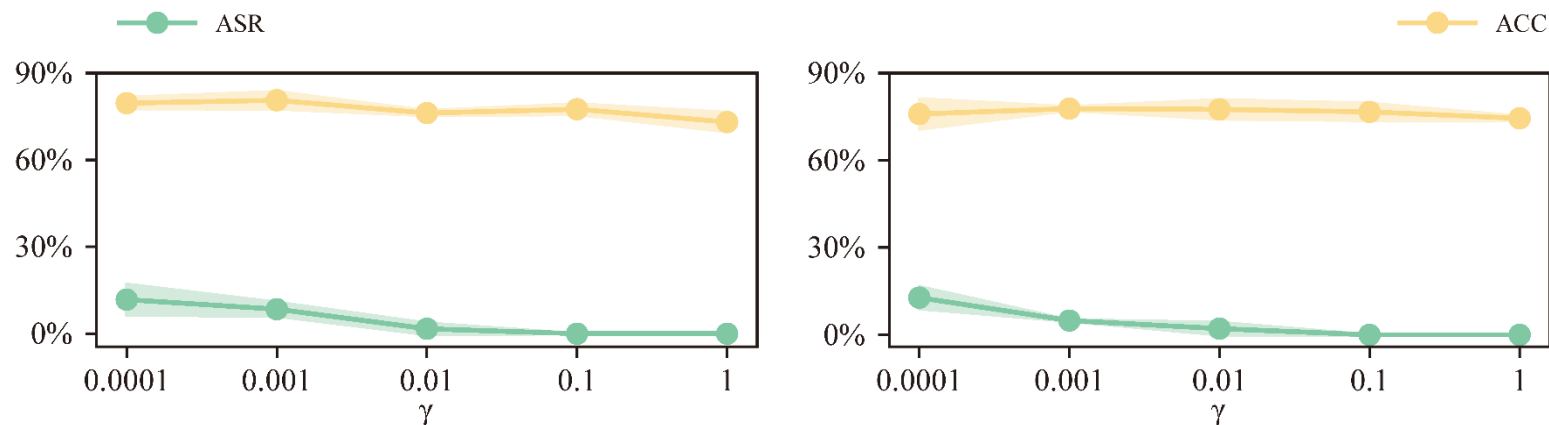
- The efficacy of DShield stems from the execution of the self-supervised learning framework and UMAP on the manipulated graph

➤ Parameter β



- The parameter β significantly affects identification precision

➤ Parameter γ



- Increasing the parameter γ results in a decline in the victim model's performance on poisoned nodes

Extension to Graph Classification Tasks

➤ Defending against Attacks on Graph Classification Tasks

Datasets	Defenses	G-SBA		G-EBA		G-GCBA	
		ASR ↓	ACC ↑	ASR ↓	ACC ↑	ASR ↓	ACC ↑
ENZYMES	<i>no-defense</i>	100.0 $\pm$ 0.00	32.67 $\pm$ 0.94	100.0 $\pm$ 0.00	27.50 $\pm$ 3.54	100.0 $\pm$ 0.00	23.33 $\pm$ 0.00
	G-DShield	01.67 $\pm$ 2.35	35.00 $\pm$ 2.36	02.50 $\pm$ 3.54	31.67 $\pm$ 2.35	00.00 $\pm$ 0.00	30.84 $\pm$ 1.18
PROTEINS	<i>no-defense</i>	100.0 $\pm$ 0.00	66.67 $\pm$ 1.27	99.56 $\pm$ 0.63	71.62 $\pm$ 3.18	99.11 $\pm$ 0.00	73.43 $\pm$ 5.73
	G-DShield	00.00 $\pm$ 0.00	74.63 $\pm$ 4.04	04.29 $\pm$ 6.06	74.33 $\pm$ 3.19	00.98 $\pm$ 1.39	71.17 $\pm$ 1.27
MNIST	<i>no-defense</i>	100.0 $\pm$ 0.00	36.33 $\pm$ 0.67	100.0 $\pm$ 0.00	36.89 $\pm$ 0.02	100.0 $\pm$ 0.00	38.34 $\pm$ 0.66
	G-DShield	00.00 $\pm$ 0.00	38.86 $\pm$ 0.37	01.87 $\pm$ 2.64	39.77 $\pm$ 1.65	00.00 $\pm$ 0.00	40.57 $\pm$ 1.89

- Experimental results show that semantic drift and attribute over-emphasis also occur in graph classification tasks and the defense efficiency is not limited to node classification backdoor attacks

Thank You!

Hao Yu

csyuhao@gmail.com

National University of Defense Technology

Source Code: <https://github.com/csyuhao/DShield>