

# Memorization in Graph Neural Networks

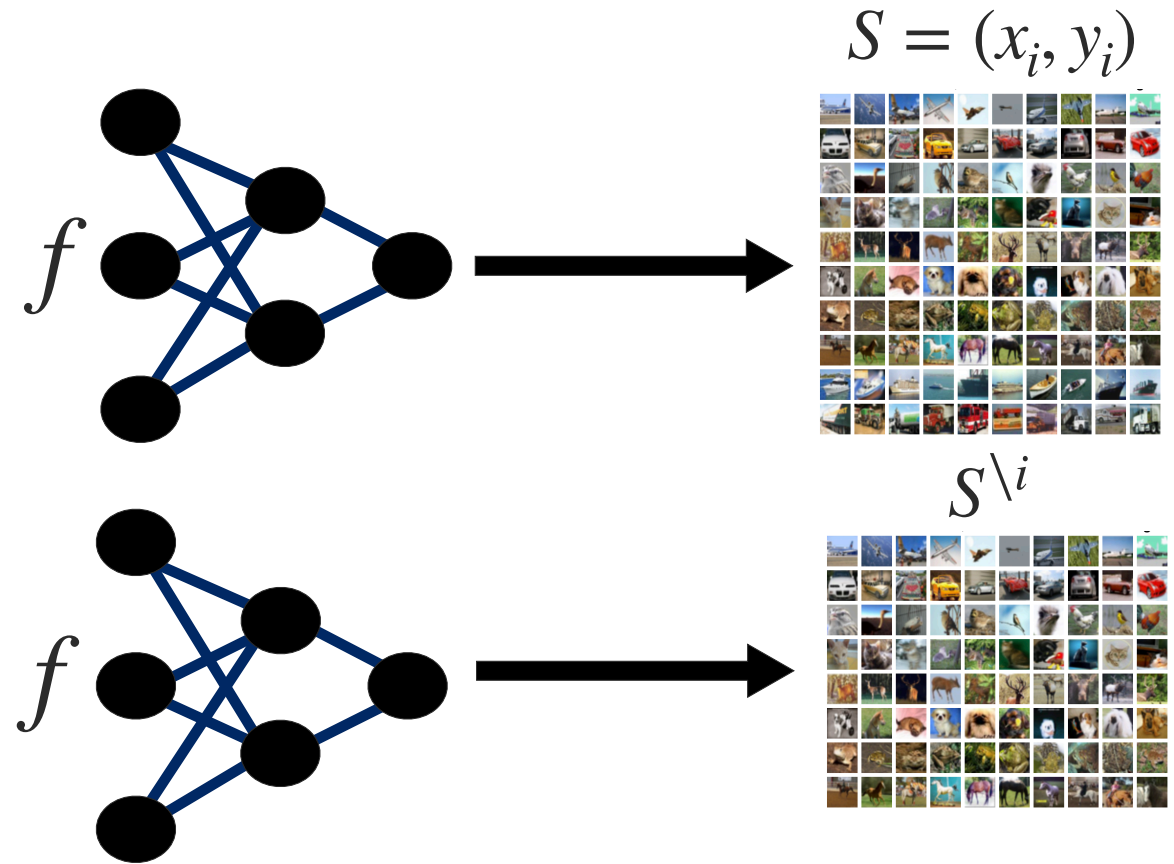
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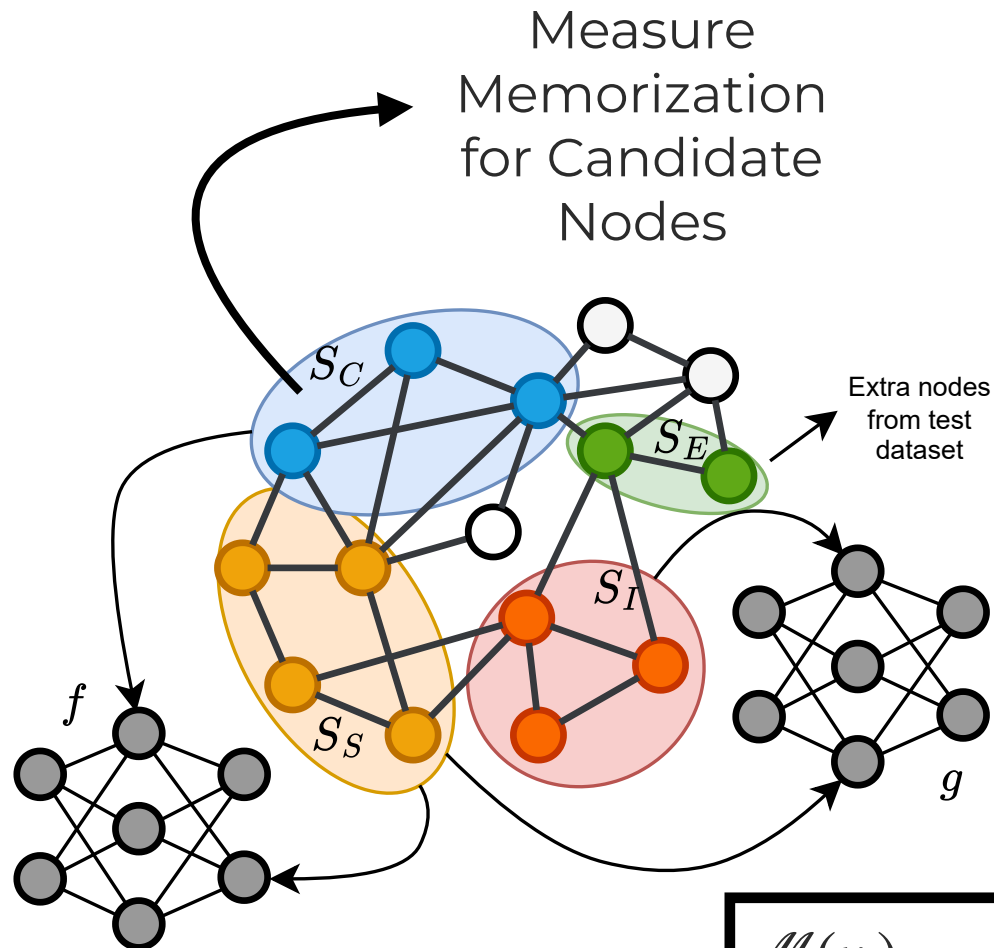
# Measuring Memorization

- Train model  $f$  on  $S$  and on  $S \setminus i$ .
- Compare the behavior of the two models.
- A model needs to see the label of the sample to correctly predict the label  $\rightarrow$  Memorized.



$$\mathcal{M}(x_i) = \mathbb{E}_{f \sim \mathcal{T}(S)} [\Pr[f(x_i) = y_i]] - \mathbb{E}_{f \sim \mathcal{T}(S \setminus x_i)} [\Pr[f(x_i) = y_i]]$$

# Measuring Memorization in GNNs



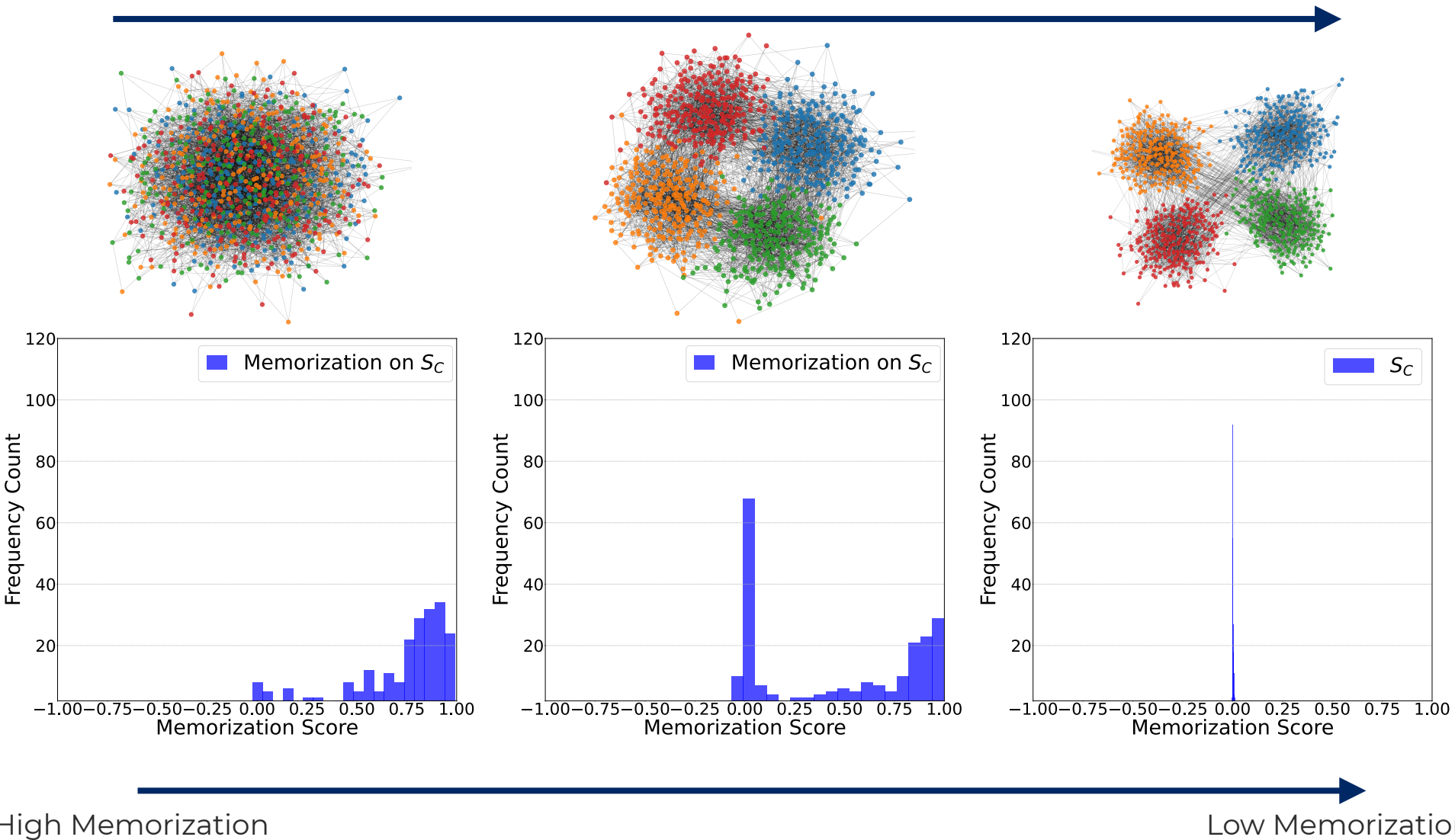
- Models  $f = S_s \cup S_c$  and  $g = S_s \cup S_I$  are trained on various subsets of nodes.
- We isolate the effect of one node's label on model behavior.

$$\mathcal{M}(v_i) = \mathbb{E}_{f \sim \mathcal{T}(S)} [\Pr[f(v_i) = y_i]] - \mathbb{E}_{g \sim \mathcal{T}(S \setminus x_i)} [\Pr[g(v_i) = y_i]]$$

# Homophily and Memorization

Low Homophily

High Homophily



# Explaining Memorization in GNNs

- We uncover 3 internal mechanisms to explain the emergence of memorization in GNNs.
- Graph homophily - like nodes connected to like nodes.
- Implicit bias of GNNs to leverage the graph structure.
- Label-Feature Inconsistency.

# Analyzing the Training Dynamics of GNNs

- We will define an alignment metric, cosine-similarity-like applied to matrices given by  $\mathcal{A}(\mathbf{K}_1, \mathbf{K}_2) = \frac{\langle \mathbf{K}_1, \mathbf{K}_2 \rangle_F}{\|\mathbf{K}_1\|_F \|\mathbf{K}_2\|_F}$
- Adjacency Matrix:  $\mathbf{A}$
- Optimal Kernel Matrix:  $\Theta^* = \bar{\mathbf{Y}}\bar{\mathbf{Y}}^T$
- NTK Matrix:  $\Theta_t^l(x, \tilde{x}; \mathbf{A}) = \nabla_W f(x; \mathbf{A})^T \cdot \nabla_W f(\tilde{x}; \mathbf{A})$

# We Will Track

## Kernel-Graph Alignment

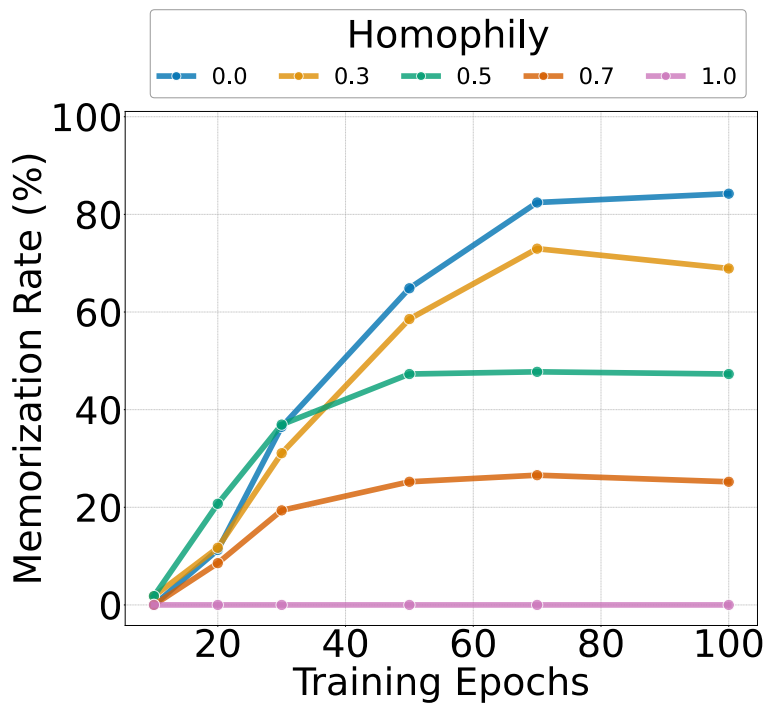
- Alignment between NTK matrix  $\Theta_t$  and adjacency matrix  $\mathbf{A}$  ( $\mathcal{A}(\Theta_t, \mathbf{A})$ ).
- Represents the **implicit bias of GNNs** to leverage the graph structure.

## Kernel-Target Alignment

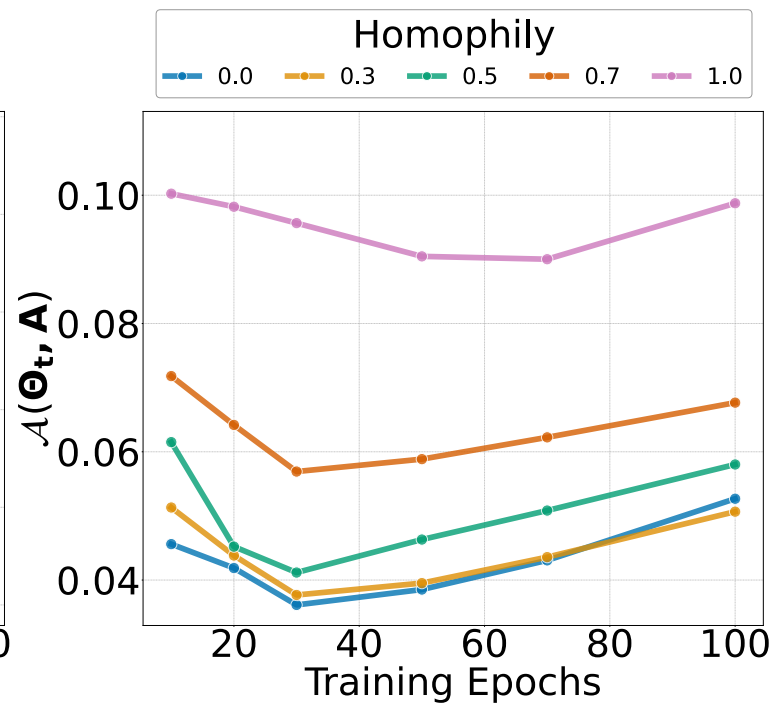
- Alignment between the NTK matrix  $\Theta_t$  and optimal kernel matrix  $\Theta^*$ .
- This metric **measures how well a classifier generalizes**, a higher alignment implies good generalization.

# Alignment Matrices for Synthetic Datasets

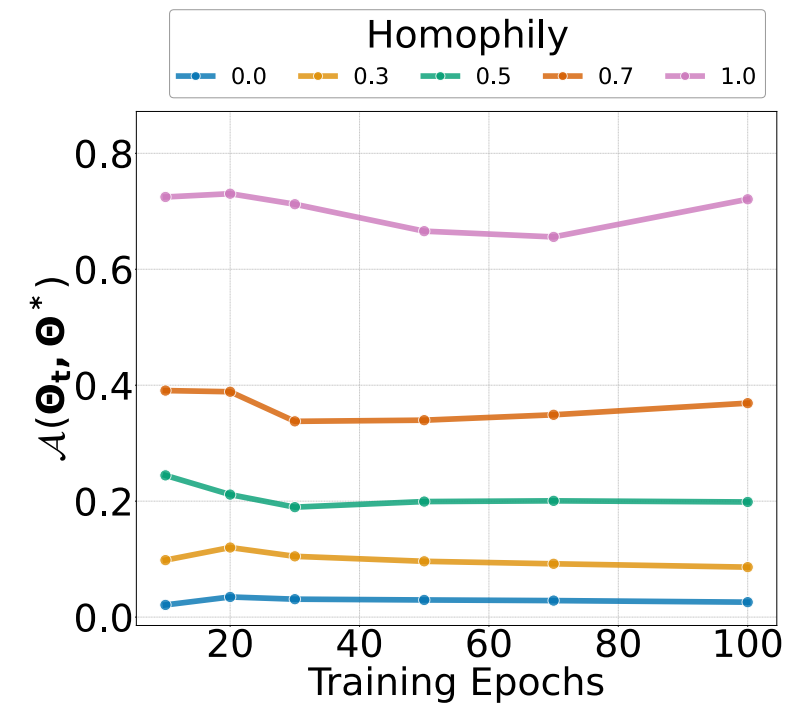
- Low homophily  $\rightarrow$  Memorization rate increases.
- Low homophily (graph structure is less informative)  $\rightarrow$  Still  $\mathcal{A}(\Theta_t, \mathbf{A})$  improves.
- Low homophily  $\rightarrow \mathcal{A}(\Theta_t, \Theta^*)$  poor, suggests memorization.



Memorization Rate



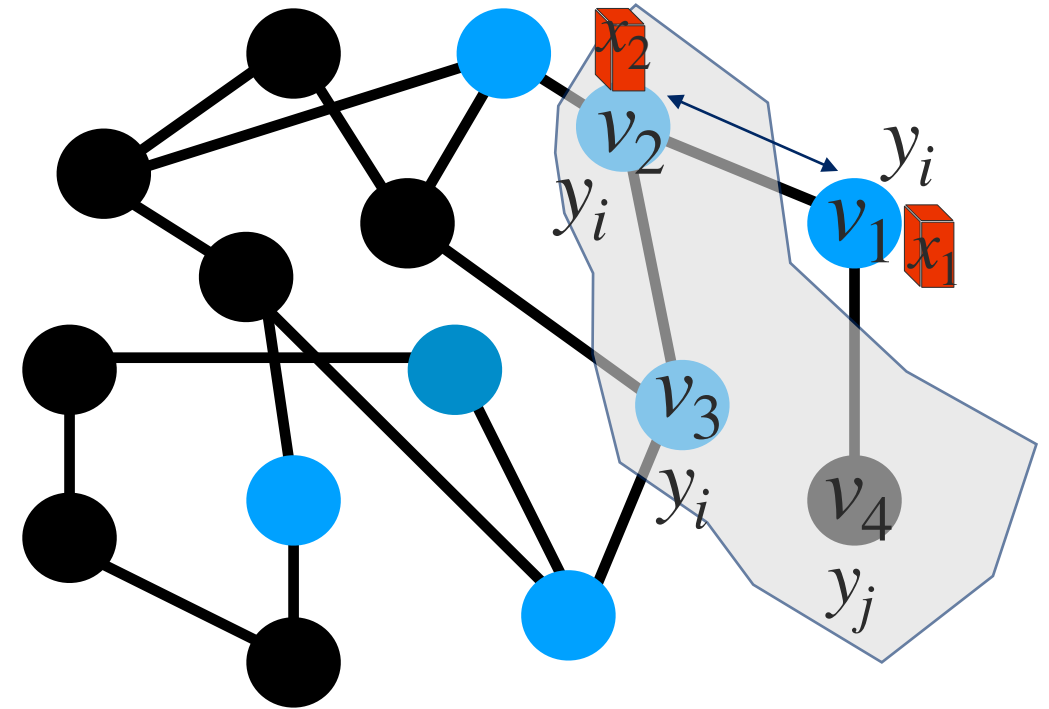
Kernel-Graph Alignment



Kernel-Target Alignment

# Node Atypicality

- Novel Label Disagreement Score (LDS) → local structural anomaly in the feature space of the nodes.
- Nodes with high LDS more likely to get memorized.



$$\text{LDS}_k(v_i) = \frac{1}{k} \sum_{v_j \in N_k(v_i)} \mathbb{I}[y_j \neq y_i]$$

# Summary

- GNNs also memorize node labels.
- Homophily $\uparrow$  Memorization Rate $\downarrow$ .
- In low-homophily settings, the graph is unhelpful for the task. But GNNs have an implicit bias to use the graph structure.
- How to achieve 0 train loss? Memorize!
- Nodes with high label disagreement score usually get memorized.

Our Paper:

