

Restructuring Graph for Higher Homophily via Adaptive Spectral Clustering

• **Abstract:** Although the ability to handle less-homophilic graphs is restricted, classical GNNs still stand out in several nice properties such as efficiency, simplicity, and explainability. In this work, we propose a novel graph restructuring method that can be integrated into any type of GNNs, including classical GNNs, to leverage the benefits of existing GNNs while alleviating their limitations. Our method learns to cluster nodes using eigenvectors beyond spectral clustering. We also proposed a new density-aware homophilic metric to better reflect the homophily of a graph.

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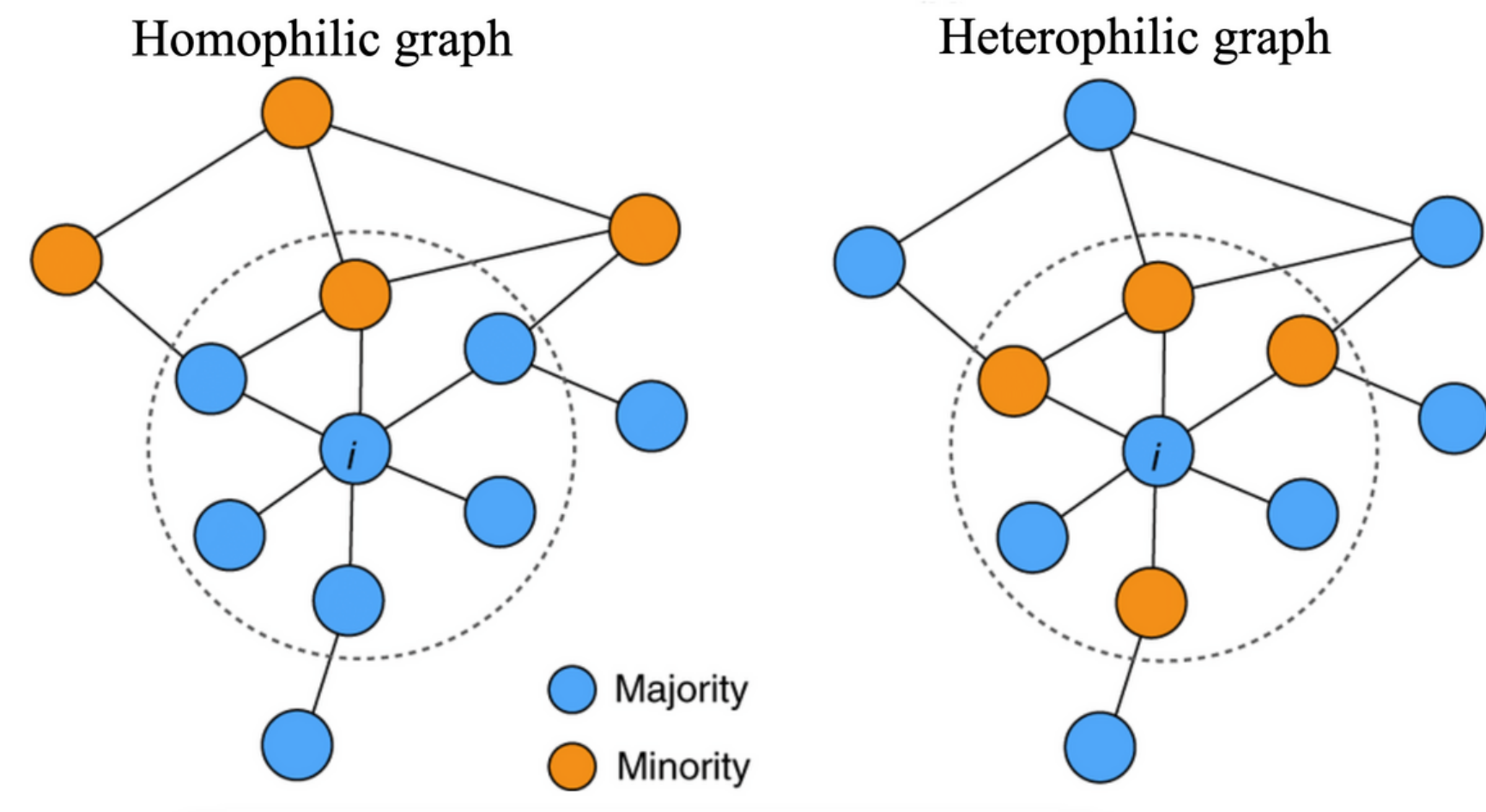
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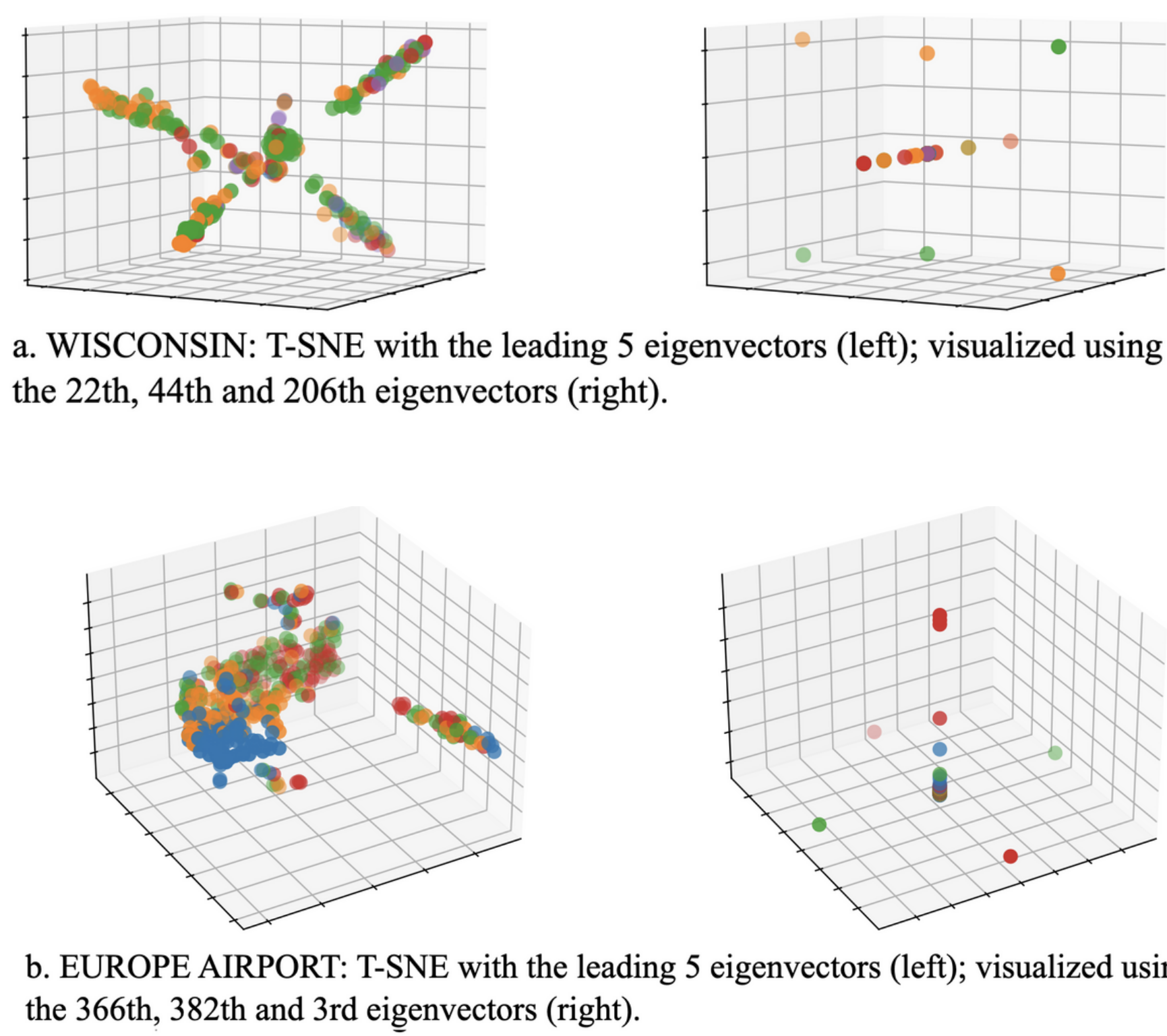
²CSE & GSAI, POSTECH

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Graphs



Spectral clustering



Density-aware homophily metric

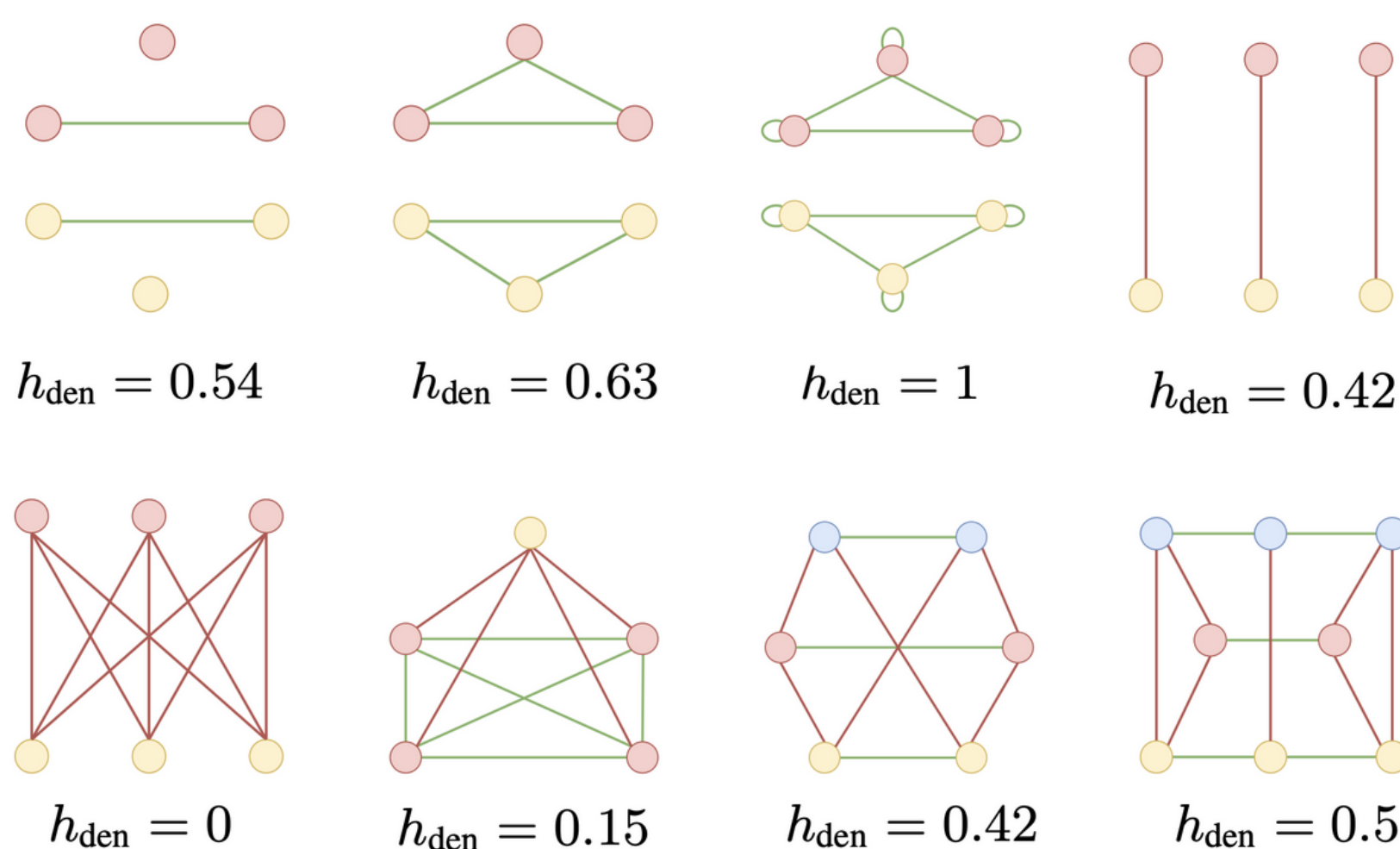
$$d_k = \frac{|(u, v) \in E : k_u = k_v = k|}{|Y_k||Y_k|},$$

$$\bar{d}_k = \max\{d_{kj} : j = 0, \dots, K-1; j \neq k\}.$$

$$\hat{h}_{\text{den}} = \min\{d_k - \bar{d}_k\}_{k=0}^{K-1}$$

↗ Intra-class edge density
↖ Inter-class edge density

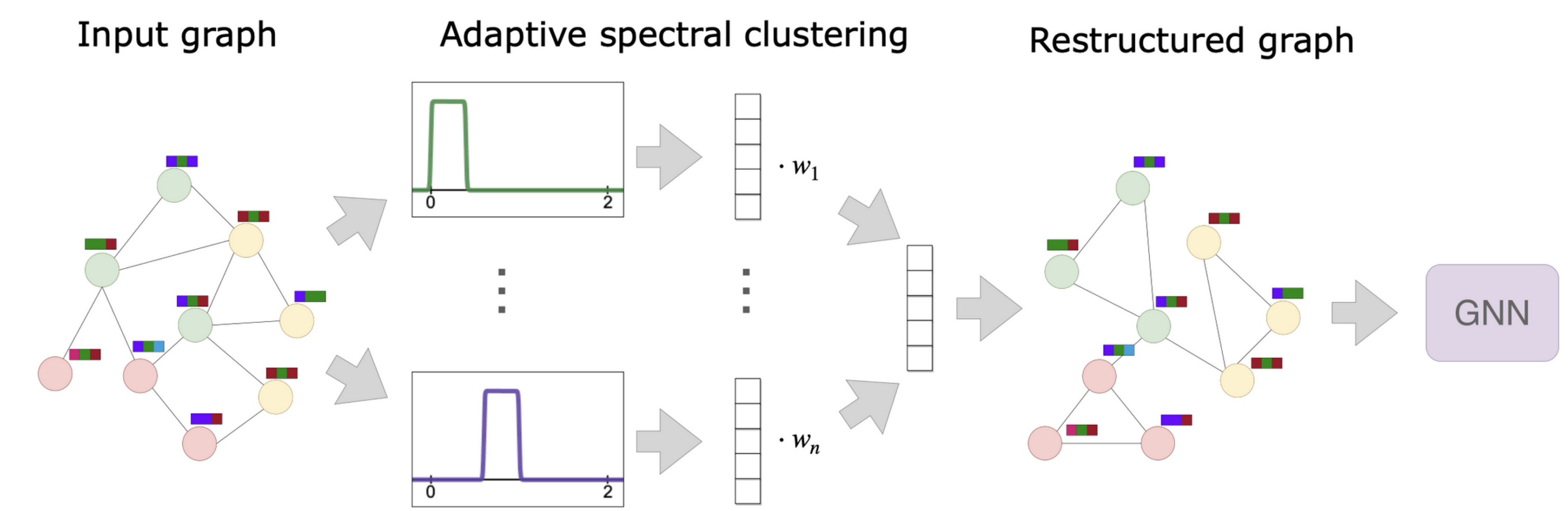
$$h_{\text{den}} = \frac{1 + \hat{h}_{\text{den}}}{2}$$



Selected references

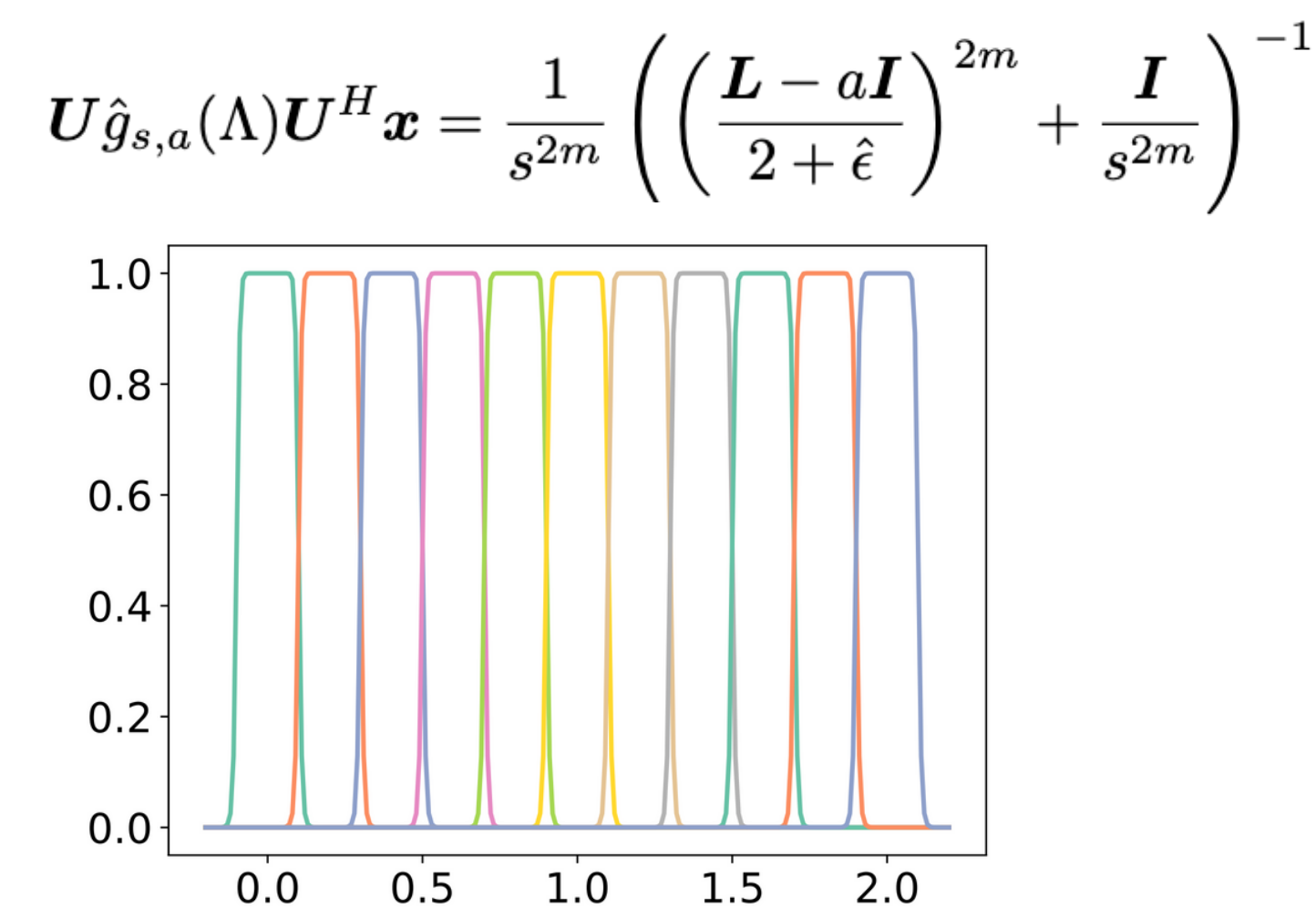
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Restructure a graph



Adaptive spectral clustering

Rectangular band-pass filters



Spectral clustering as spectral filtering

$$\Gamma_{s,a} = \hat{g}_{s,a}(L)(R \frown X)$$

Laplacian matrix
Random node feature

Rectangular band-pass filter
Random projection (Johnson-Lindenstrauss lemma)

Concatenation
Learnable function

$$\Gamma = (\Gamma_{s_1, a_1} \frown \Gamma_{s_2, a_2} \frown \dots)$$

$$H = \Theta(\Gamma).$$

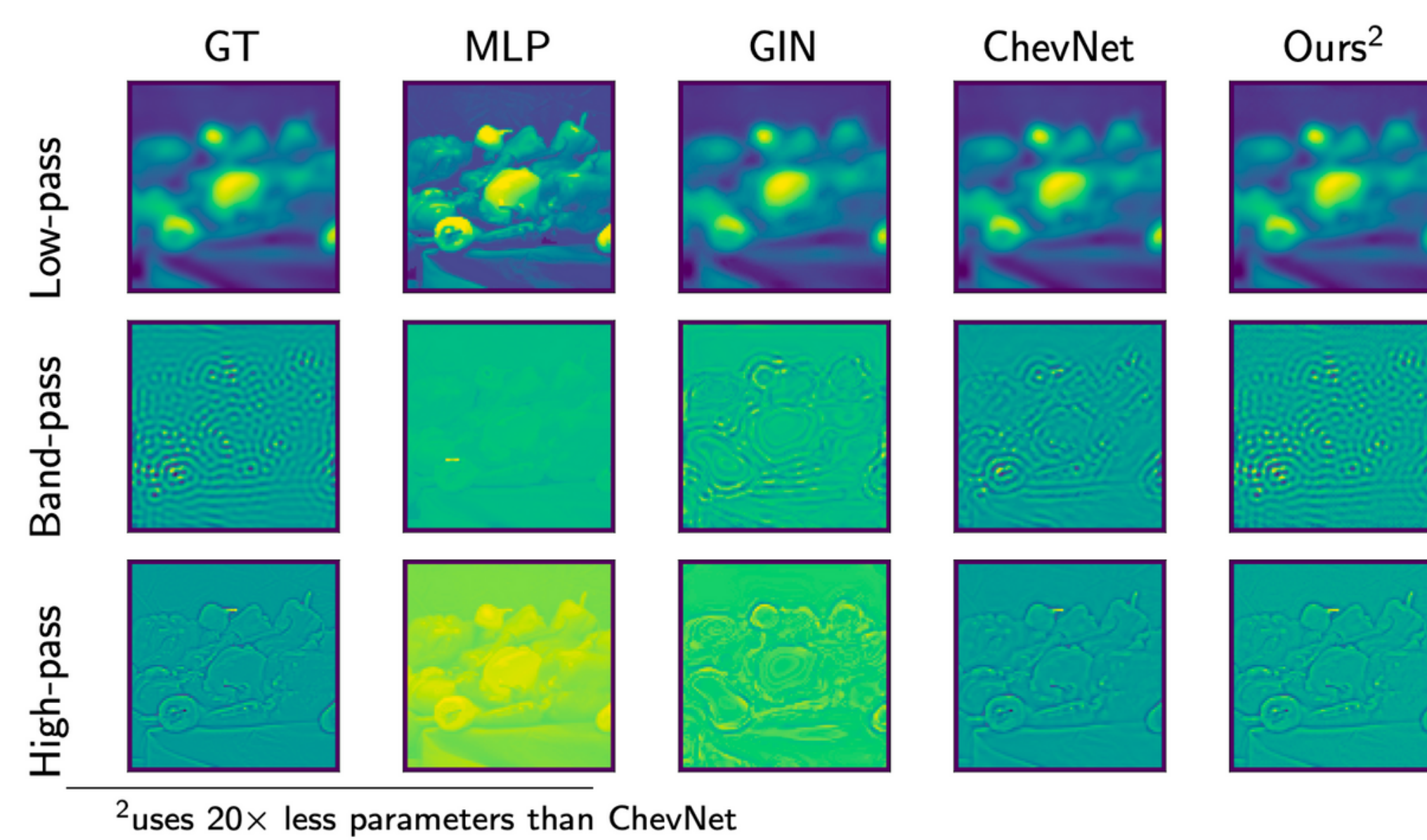
Triplet loss function

$$\mathcal{L}(\Theta) = \sum_{\substack{i,j \in V_Y \\ k \in N_Y(i) \\ y_i = y_j}} [||H_{i \cdot} - H_{j \cdot}||^2 - ||H_{i \cdot} - H_{k \cdot}||^2 + \epsilon]_+$$

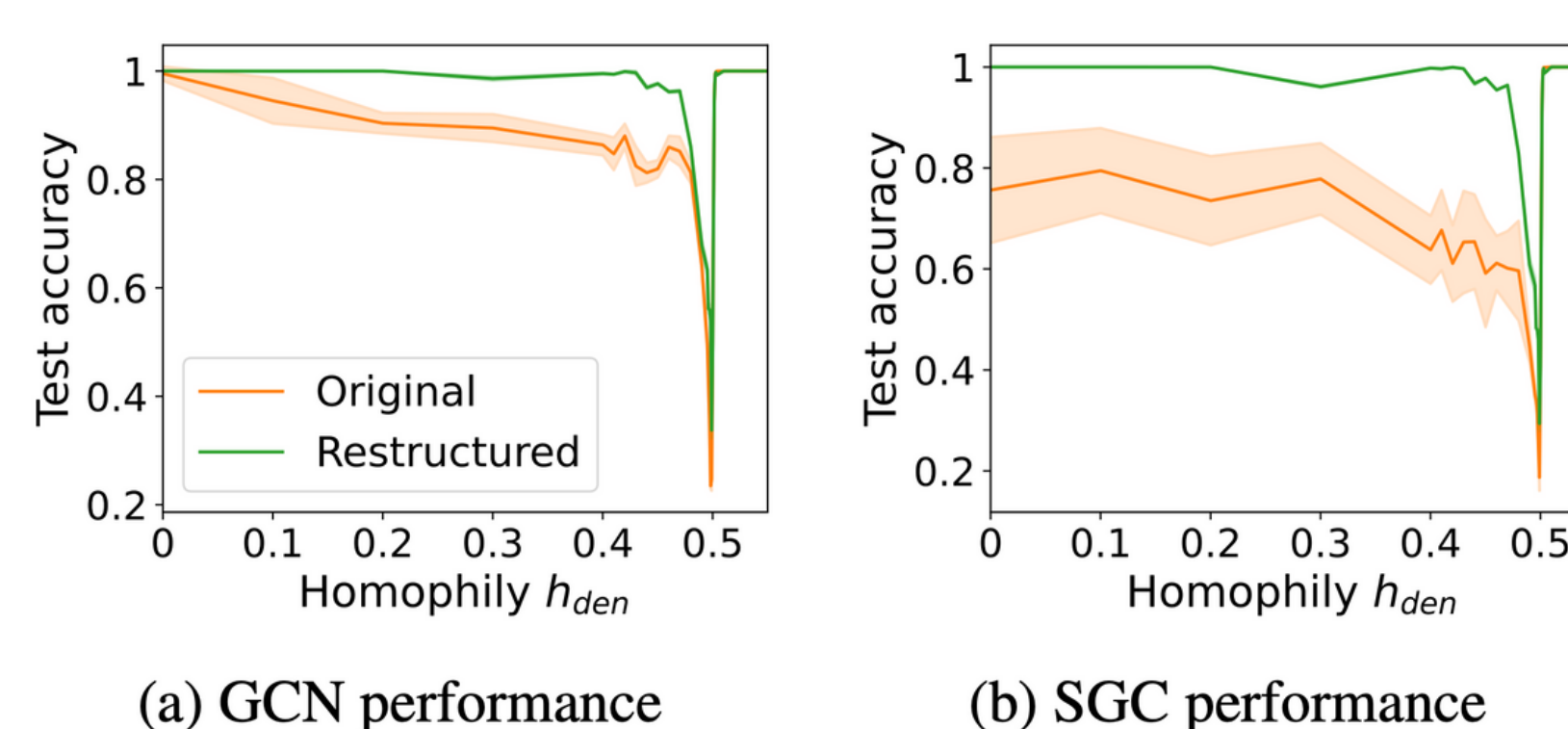
positive pairs
negative pairs

Experimental results

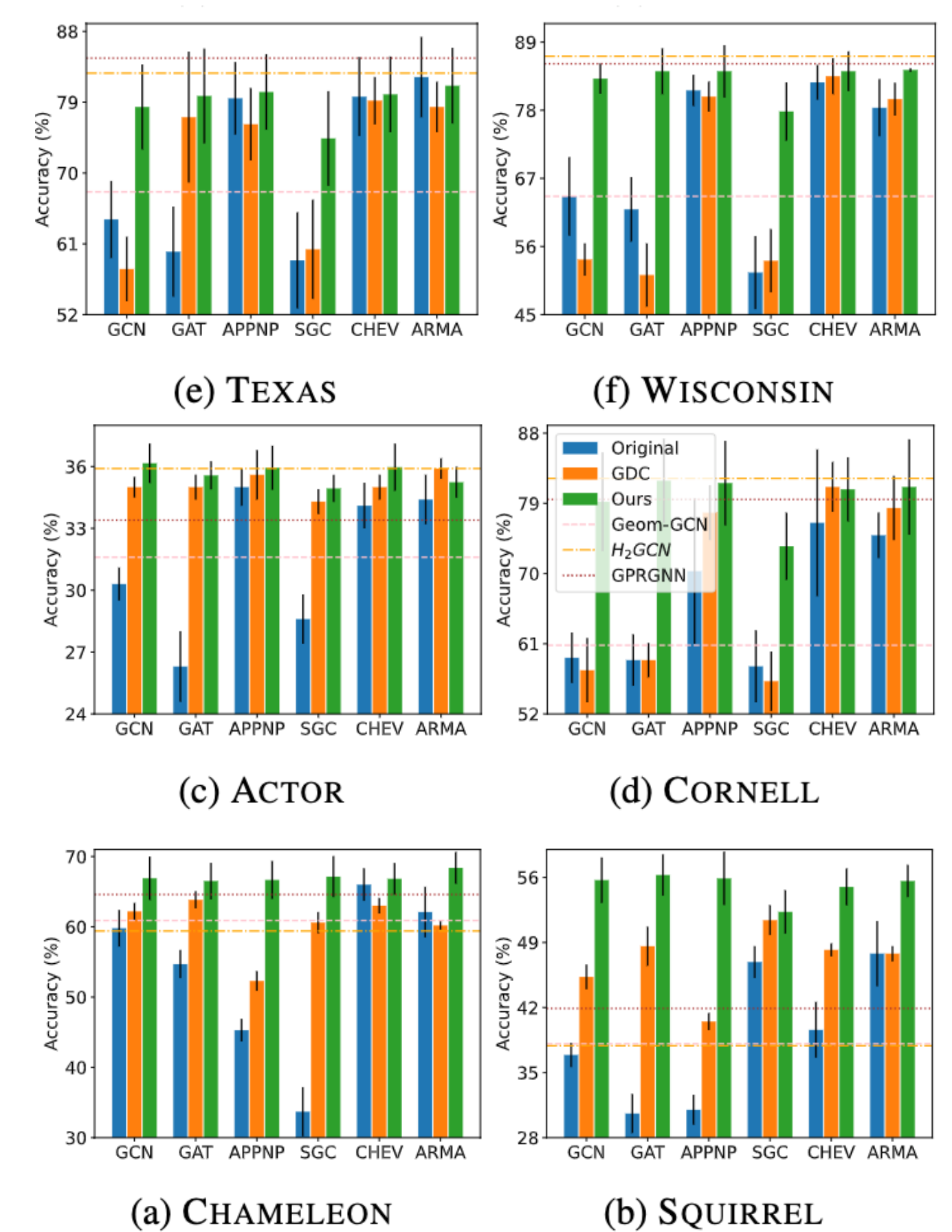
Spectral expressivity



Synthetic heterophilic graphs



Node classification



- The model can best recover images filtered using low-, band- and high-pass functions.
- The restructured real-world graphs have higher homophily which improves GNNs performance by an average 25%.
- The method yields high accuracy on synthetic heterophilic graphs.