

Less is More: SlimG for Accurate, Robust, and Interpretable Graph Mining



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Outline



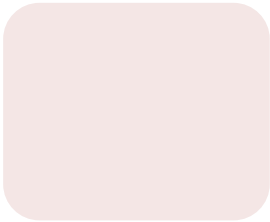
- Problem Definition
- Q1: Reasons
- Q2: Method
- Q3: Sanity Checks
- Experiments
- Conclusions

Outline

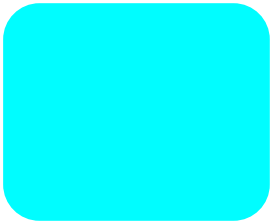


- ➔ • Problem Definition
 - Motivation
 - Research Questions
- Q1: Reasons
- Q2: Method
- Q3: Sanity Checks
- Experiments
- Conclusions

Legend



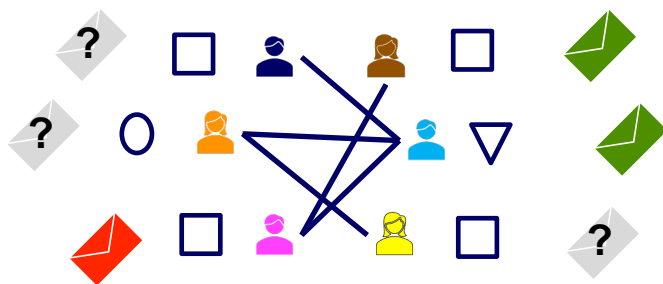
→ Outline/Roadmap



→ Reminder



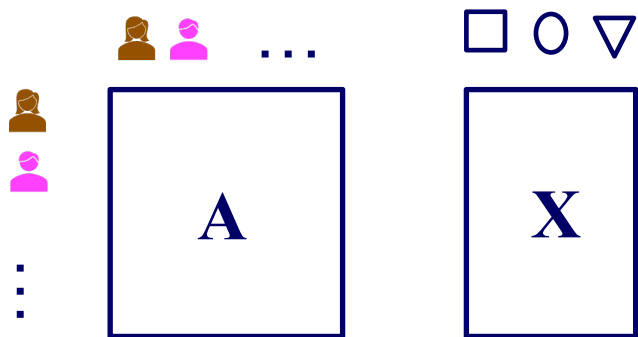
Task Definition



Given:

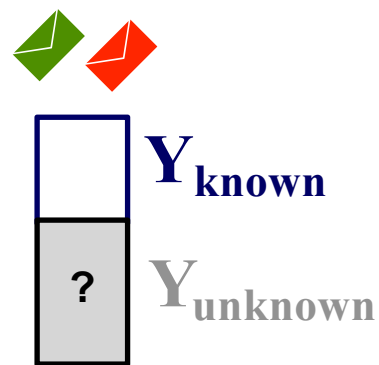
- Graph A
- Node features X
- A few node labels

Predict: Rest labels



Adjacency
Matrix

Node
Features

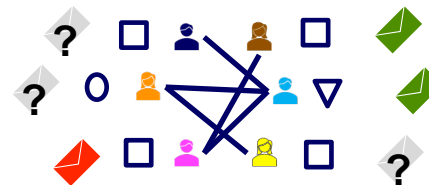


Labels

Task Definition

- Given
 - Graph (single edge-type)
 - Node features (black or green hair, etc.)
 - A few class labels ('vote', e.g., R/G)

- Predict
 - Rest labels



OUR Problem Definition

- Find good ‘derived features’ \equiv embeddings for each node
 - (Using A , X , AX , etc.)
- That best predict the unknown labels $Y_{unknown}$

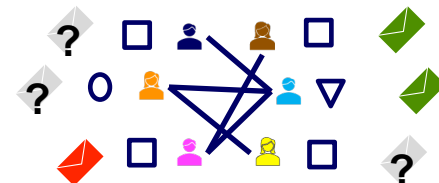


Graph Neural Networks (GNNs)

- GNNs have shown their ability to handle different tasks in graph domain:
 - GCN
 - GAT
 - ...
- Some GNNs remove the non-linear functions in their models:
 - SGC
 - DGC
 - ...

Research Questions

- Q1) Reasons
- Q2) GNN that is
 - Accurate, Robust, Fast, Scalable, Interpretable
- Q3) Sanity Checks



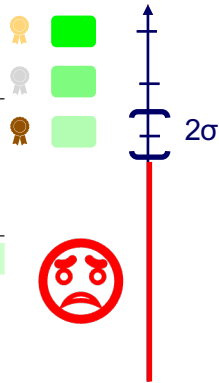
Sneak Preview: Accurate & Robust

Real-World Datasets

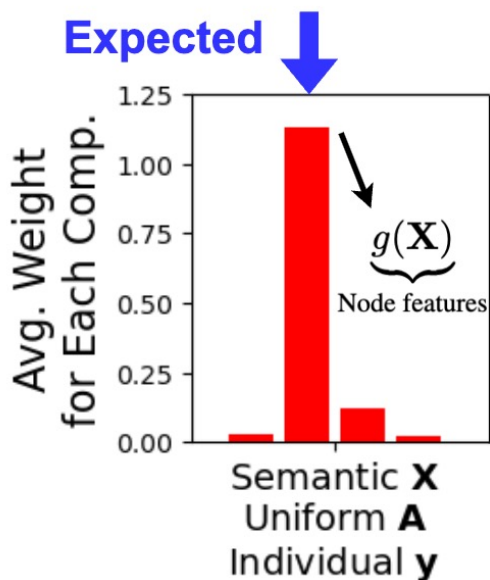
7 Homophily

6 Heterophily

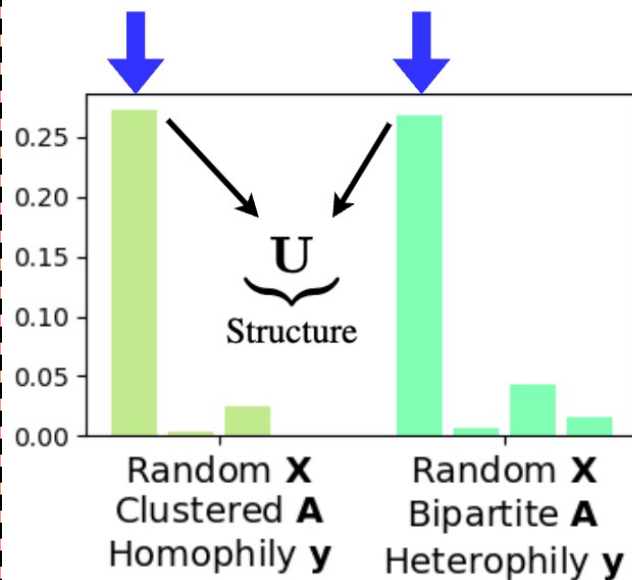
Model	Cora	CiteSeer	PubMed	Comp.	Photo	ArXiv	Products	Cham.	Squirrel	Actor	Penn94	Twitch	Pokec	Avg. Rank
LR	51.5±1.2	52.9±4.5	79.9±0.5	73.9±1.2	79.3±1.5	48.3±1.9	56.4±0.5	24.9±1.7	26.7±1.9	27.8±0.8	63.5±0.5	53.0±0.1	61.3±0.0	11.7 (4.2)
Reg. Kernel	67.8±2.5	62.1±4.4	83.4±1.4	80.3±1.4	87.1±1.2	O.O.M.	O.O.M.	29.4±2.6	24.3±2.3	29.6±1.4	O.O.M.	O.O.M.	O.O.M.	12.2 (3.8)
Diff. Kernel	70.6±1.5	62.7±3.8	82.1±0.4	83.1±1.0	89.8±0.6	O.O.M.	O.O.M.	34.5±7.9	28.3±1.5	24.7±0.9	53.5±0.8	O.O.M.	O.O.M.	11.8 (2.5)
RW Kernel	72.7±1.7	64.1±3.9	83.1±0.7	84.2±0.7	90.6±0.7	63.2±0.2	74.2±0.0	34.9±3.5	25.0±1.6	26.4±1.1	63.1±0.7	57.6±0.1	59.5±0.0	8.3 (3.3)
SGC	76.2±1.1	65.8±3.9	84.1±0.8	83.7±1.6	90.1±0.9	65.0±3.4	74.6±5.1	38.1±4.5	33.1±1.0	24.6±0.8	64.0±1.1	56.5±0.1	69.8±0.0	6.6 (4.2)
DGC	77.8±1.4	66.1±4.2	84.3±0.6	83.9±0.7	90.4±0.2	65.2±4.0	68.7±13.	37.2±3.7	29.2±1.2	25.2±2.1	62.5±0.4	58.2±0.2	60.7±0.1	6.6 (3.2)
S ² GC	78.3±1.5	66.9±4.4	84.3±0.3	83.1±0.8	90.1±0.8	62.0±7.4	58.3±18.	34.9±4.9	27.6±1.8	26.7±1.8	63.1±0.5	58.7±0.1	61.2±0.0	6.6 (2.7)
G ² CN	76.6±1.5	64.2±3.3	81.4±0.6	82.8±1.6	88.8±0.5	O.O.M.	O.O.M.	40.7±2.9	32.1±1.5	24.3±0.5	O.O.M.	O.O.M.	O.O.M.	10.5 (4.5)
GCN	76.0±1.2	65.0±2.9	84.3±0.5	85.1±0.9	91.6±0.5	62.8±0.6	O.O.M.	38.5±3.0	31.4±1.8	26.8±0.4	62.9±0.7	57.0±0.1	63.9±0.4	6.3 (2.4)
SAGE	74.6±1.3	63.7±3.6	82.9±0.4	83.8±0.5	90.6±0.5	61.5±0.6	O.O.M.	39.8±4.3	27.0±1.3	27.8±0.9	O.O.M.	56.6±0.4	68.9±0.1	8.5 (3.5)
GCNII	77.8±1.7	63.4±3.0	84.9±0.8	82.3±1.8	90.8±0.6	45.7±0.5	O.O.M.	30.5±2.5	21.9±3.0	29.0±1.3	64.5±0.5	56.9±0.6	62.1±0.3	8.4 (4.6)
H ² GCN	77.6±0.9	64.7±3.8	85.4±0.4	49.5±16.	75.8±11.	O.O.M.	O.O.M.	31.9±2.6	25.0±0.5	28.9±0.6	63.9±0.4	58.7±0.0	O.O.M.	8.9 (4.9)
APPNP	80.0±0.6	67.1±2.8	84.6±0.5	84.2±1.7	92.5±0.3	53.4±1.3	O.O.M.	30.9±4.7	23.9±3.2	26.1±1.0	63.7±0.9	47.3±0.3	57.4±0.4	7.6 (4.8)
GPR-GNN	78.8±1.3	64.2±4.0	85.1±0.7	85.0±1.0	92.6±0.3	58.5±0.8	O.O.M.	31.7±4.7	26.2±1.6	29.5±1.1	64.5±0.4	57.6±0.2	67.6±0.1	5.4 (3.7)
GAT	78.2±1.2	65.8±4.0	83.6±0.2	85.4±1.4	91.7±0.5	58.2±1.0	O.O.M.	39.1±4.1	28.6±0.6	26.4±0.4	60.5±0.8	O.O.M.	O.O.M.	7.5 (3.7)
SLIMG	77.8±1.1	67.1±2.3	84.6±0.5	86.3±0.7	91.8±0.5	66.3±0.3	84.9±0.0	40.8±3.2	31.1±0.7	30.9±0.6	68.2±0.6	59.7±0.1	73.9±0.1	1.9 (1.5)



Sneak Preview: Interpretable



(a) No network effects

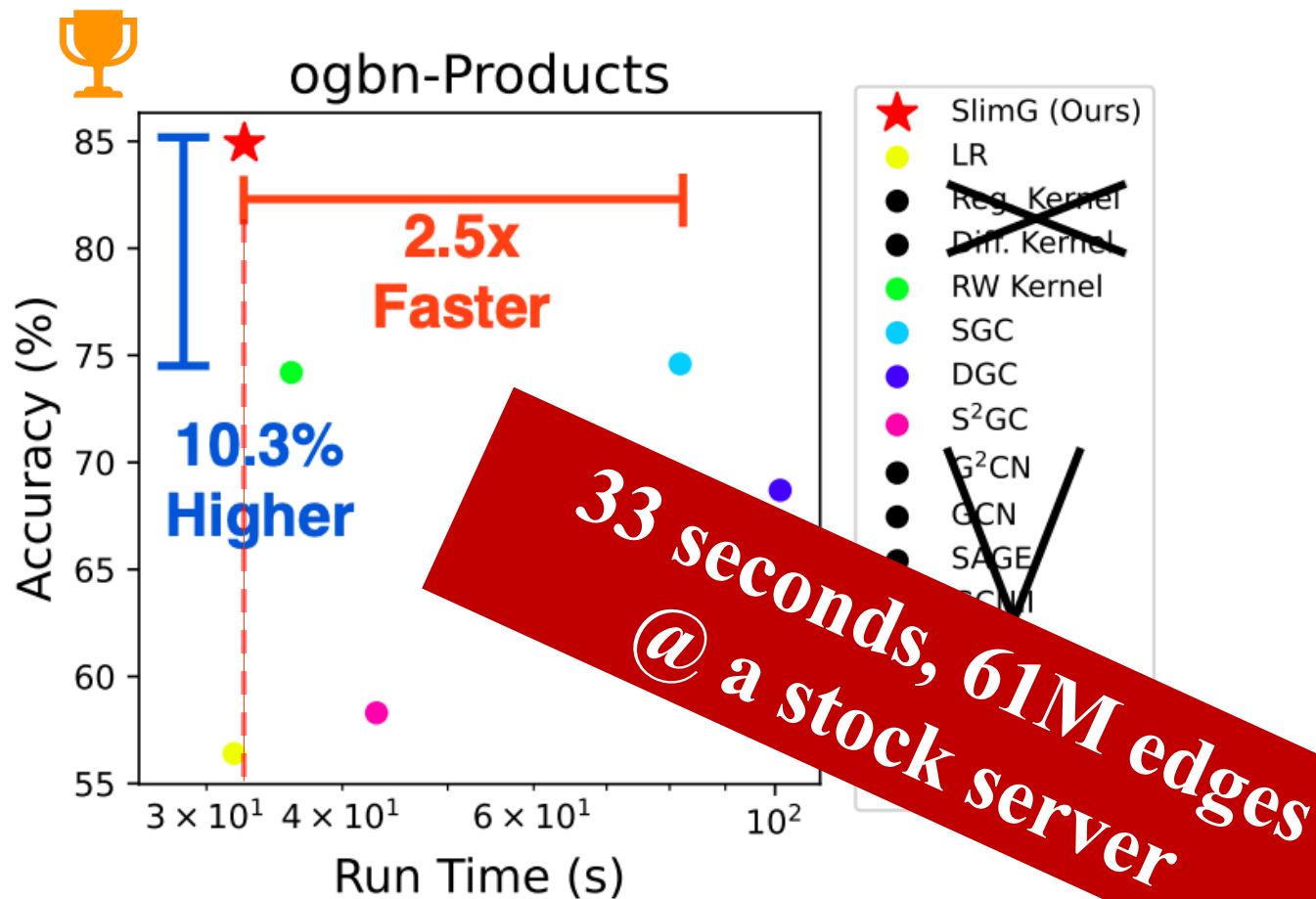


(b) Useless features

$$\underbrace{\mathbf{U}}_{\text{Structure}} \parallel \underbrace{g(\mathbf{X})}_{\text{Node features}} \parallel \underbrace{g(\mathbf{A}_{\text{row}}^2 \mathbf{X})}_{\text{2-step neighbors}} \parallel \underbrace{g(\tilde{\mathbf{A}}_{\text{sym}}^2 \mathbf{X})}_{\text{Neighbors}}$$

$$\underbrace{\mathbf{U}}_{\text{Structure}} \parallel \underbrace{g(\mathbf{X})}_{\text{Node features}} \parallel \underbrace{g(\mathbf{A}_{\text{row}}^2 \mathbf{X})}_{\text{2-step neighbors}} \parallel \underbrace{g(\tilde{\mathbf{A}}_{\text{sym}}^2 \mathbf{X})}_{\text{Neighbors}}$$

Sneak Preview: Fast & Scalable



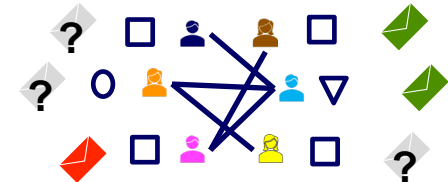
Research Questions

- Q1) Reasons

- Q2) Goal **“Careful Simplicity”**

– Accurate, Robust, Fast, Scalable, Interpretable

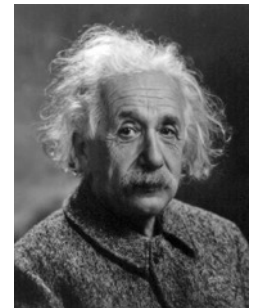
- Q3) Sanity Checks



Over-Arching Principle

“Careful Simplicity”

“Everything should be made as simple as possible, but not simpler.”



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Outline



- Problem Definition
- ➔ • Q1: Reasons
- Q2: Method
- Q3: Sanity Checks
- Experiments
- Conclusions

Q1: Reasons – Why?

- Why/when do GNNs succeed,
- ... and why/when do they fail?

- Q: How should we proceed?



Model

LR

SGC

DGC

S^2GC

G^2CN

PPNP*

APPNP*

GDC*

GPR-GNN*

ChebNet*

GCN*

SAGE*

GCNII*

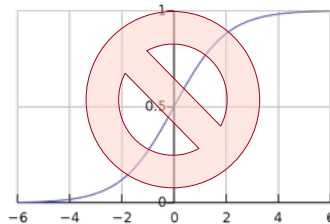
H_2GCN^*

GAT**

DA-GNN**

Q1: Reasons – Why?

- Why/when do GNNs succeed,
- ... and why/when do they fail?
- Q: How should we proceed?
- A: Drop non-linearities, to get the essence





GNNEXP



Jaemin Yoo

Model	Type	Propagator function $\mathcal{P}(\mathbf{A}, \mathbf{X})$
LR	Linear	\mathbf{X}
SGC	Linear	$\tilde{\mathbf{A}}_{\text{sym}}^K \mathbf{X}$
DGC	Linear	$[(1 - T/K)\mathbf{I} + (T/K)\tilde{\mathbf{A}}_{\text{sym}}]^K \mathbf{X}$
S ² GC	Linear	$\sum_{k=1}^K (\alpha\mathbf{I} + (1 - \alpha)\tilde{\mathbf{A}}_{\text{sym}}^k) \mathbf{X}$
G ² CN	Linear	$\prod_{i=1}^N [\mathbf{I} - (T_i/K)((b_i - 1)\mathbf{I} + \mathbf{A}_{\text{sym}})^2]^K \mathbf{X}$
PPNP*	Decoupled	$(\mathbf{I} - (1 - \alpha)\tilde{\mathbf{A}}_{\text{sym}})^{-1} \mathbf{X}$
APPNP*	Decoupled	$[\sum_{k=0}^{K-1} \alpha(1 - \alpha)^k \tilde{\mathbf{A}}_{\text{sym}}^k + (1 - \alpha)^K \tilde{\mathbf{A}}_{\text{sym}}^K] \mathbf{X}$
GDC*	Decoupled	$\mathbf{S} = \text{sparse}_{\epsilon}(\sum_{k=0}^{\infty} (1 - \alpha)^k \tilde{\mathbf{A}}_{\text{sym}}^k)$ for $\tilde{\mathbf{S}}_{\text{sym}} \mathbf{X}$
GPR-GNN*	Decoupled	$\prod_{k=0}^K \tilde{\mathbf{A}}_{\text{sym}}^k \mathbf{X}$
ChebNet*	Coupled	$\prod_{k=0}^{K-1} \mathbf{A}_{\text{sym}}^k \mathbf{X}$
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SAGE*	Coupled	$\prod_{k=0}^K \mathbf{A}_{\text{sym}}^{\text{row} k} \mathbf{X}$
GCNII*	Coupled	$\prod_{k=0}^{K-2} \tilde{\mathbf{A}}_{\text{sym}}^k \mathbf{X} \parallel ((1 - \alpha)\tilde{\mathbf{A}}_{\text{sym}}^K + \alpha\tilde{\mathbf{A}}_{\text{sym}}^{K-1}) \mathbf{X}$
H ₂ GCN*	Coupled	$\prod_{k=0}^{2K} \mathbf{A}_{\text{sym}}^k \mathbf{X}$
GAT**	Attention	$\prod_{k=1}^K [\text{diag}(\mathbf{X}\mathbf{w}_{k,1})\tilde{\mathbf{A}} + \tilde{\mathbf{A}}\text{diag}(\mathbf{X}\mathbf{w}_{k,2})] \mathbf{X}$
DA-GNN**	Attention	$\sum_{k=0}^K \text{diag}(\tilde{\mathbf{A}}_{\text{sym}}^k \mathbf{X}\mathbf{w})\tilde{\mathbf{A}}_{\text{sym}}^k \mathbf{X}$



GNNEXP



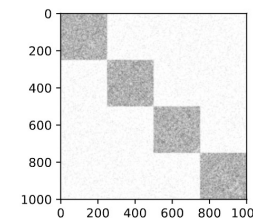
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S ² GC	Linear	$\sum_{k=1}^K (\alpha\mathbf{I} + (1 - \alpha)\tilde{\mathbf{A}}_{\text{sym}}^k) \mathbf{X}$
G ² CN	Linear	$\prod_{i=1}^N [\mathbf{I} - (T_i/K)((b_i - 1)\mathbf{I} + \mathbf{A}_{\text{sym}})^2]^K \mathbf{X}$
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DA-GNN**	Attention	$\sum_{k=0}^K \text{diag}(\tilde{\mathbf{A}}_{\text{sym}}^k \mathbf{X}\mathbf{w})\tilde{\mathbf{A}}_{\text{sym}}^k \mathbf{X}$

Q1: Reasons – Deep Dive

- Successes
 - S1: Homophily
 - S2: [Heterophily]
- Pain Points
 - PP1: Lack of Robustness
 - PP2: Vulnerability to Noisy Features
 - PP3: Efficiency and Effectiveness
 - PP4: Many Hyperparameters

Q1: Reasons – Deep Dive



- Successes
 - ✓ S1: Homophily $AX; A^2X; \dots \rightarrow$ neighbors' features
 - S2: [Heterophily]
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Q1: Reasons – Deep Dive

- Successes

✓ S1: Homophily $AX; A^2X; \dots \rightarrow$ neighbors' features

✗ ✓ S2: [Heterophily] AX : HURTS

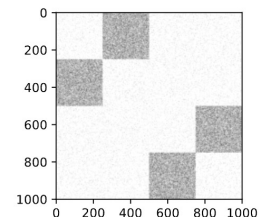
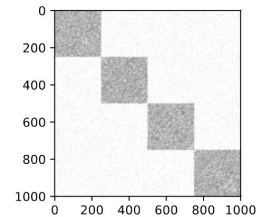
• Pain Points A^2X : if concat., good

- PP1: Lack of Robustness

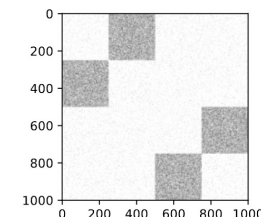
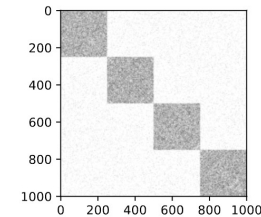
- PP2: Vulnerability to Noisy Features

- PP3: Efficiency and Effectiveness

- PP4: Many Hyperparameters



Q1: Reasons – Deep Dive



- Successes

✓ S1: Homophily $AX; A^2X; \dots \rightarrow$ neighbors' features

✗ ✓ S2: [Heterophily] AX : HURTS

A^2X : if concat., good

- Pain Points

✗ PP1: Lack of Robustness

$(\dots)X$



✗ PP2: Vulnerability to Noisy Features $(\dots + I)X$

– PP3: Efficiency and Effectiveness

– PP4: Many Hyperparameters

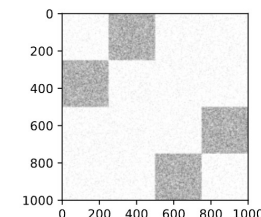
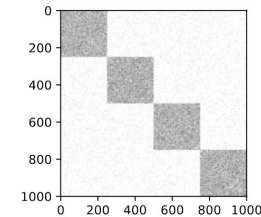
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GCNII*	Coupled	$\prod_{k=0}^{K-2} \tilde{\mathbf{A}}_{\text{sym}}^k \mathbf{X} \parallel ((1 - \alpha)\tilde{\mathbf{A}}_{\text{sym}}^K + \alpha\tilde{\mathbf{A}}_{\text{sym}}^{K-1}) \mathbf{X}$
H ₂ GCN*	Coupled	$\prod_{k=0}^{2K} \mathbf{A}_{\text{sym}}^k \mathbf{X}$
GAT**	Attention	$\prod_{k=1}^K [\text{diag}(\mathbf{X}\mathbf{w}_{k,1})\tilde{\mathbf{A}} + \tilde{\mathbf{A}}\text{diag}(\mathbf{X}\mathbf{w}_{k,2})] \mathbf{X}$
DA-GNN**	Attention	$\sum_{k=0}^K \text{diag}(\tilde{\mathbf{A}}_{\text{sym}}^k \mathbf{X}\mathbf{w})\tilde{\mathbf{A}}_{\text{sym}}^k \mathbf{X}$

Q1: Reasons – Deep Dive



- Successes

✓ S1: Homophily $AX; A^2X; \dots \rightarrow$ neighbors' features

✗ ✓ S2: [Heterophily] AX : HURTS

A^2X : if concat., good

- Pain Points

✗ PP1: Lack of Robustness $(\dots)X$ 

✗ PP2: Vulnerability to Noisy Features $(\dots + I)X$ 

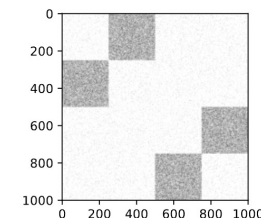
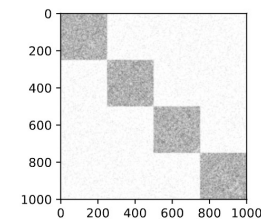
✗ PP3: Efficiency and Effectiveness 'too many cooks'

– PP4: Many Hyperparameters 

GNNEXP

Model	Type	Propagator function $\mathcal{P}(\mathbf{A}, \mathbf{X})$
LR	Linear	\mathbf{X}
SGC	Linear	$\tilde{\mathbf{A}}_{\text{sym}}^K \mathbf{X}$
DGC	Linear	$[(1 - T/K)\mathbf{I} + (T/K)\tilde{\mathbf{A}}_{\text{sym}}]^K \mathbf{X}$
S ² GC	Linear	$\sum_{k=1}^K (\alpha\mathbf{I} + (1 - \alpha)\tilde{\mathbf{A}}_{\text{sym}}^k) \mathbf{X}$
G ² CN	Linear	$\prod_{i=1}^N [\mathbf{I} - (T_i/K)((b_i - 1)\mathbf{I} + \mathbf{A}_{\text{sym}})^2]^K \mathbf{X}$
PPNP*	Decoupled	$(\mathbf{I} - (1 - \alpha)\tilde{\mathbf{A}}_{\text{sym}})^{-1} \mathbf{X}$
APPNP*	Decoupled	$[\sum_{k=0}^{K-1} \alpha(1 - \alpha)^k \tilde{\mathbf{A}}_{\text{sym}}^k + (1 - \alpha)^K \tilde{\mathbf{A}}_{\text{sym}}^K] \mathbf{X}$
GDC*	Decoupled	$\mathbf{S} = \text{sparse}_{\epsilon}(\sum_{k=0}^{\infty} (1 - \alpha)^k \tilde{\mathbf{A}}_{\text{sym}}^k)$ for $\tilde{\mathbf{S}}_{\text{sym}} \mathbf{X}$
GPR-GNN*	Decoupled	$\prod_{k=0}^K \tilde{\mathbf{A}}_{\text{sym}}^k \mathbf{X}$
ChebNet*	Coupled	$\prod_{k=0}^{K-1} \mathbf{A}_{\text{sym}}^k \mathbf{X}$
GCN*	Coupled	$\tilde{\mathbf{A}}_{\text{sym}}^K \mathbf{X}$
SAGE*	Coupled	$\prod_{k=0}^K \mathbf{A}_{\text{sym}}^k \mathbf{X}$
GCNII*	Coupled	$\prod_{k=0}^{K-2} \tilde{\mathbf{A}}_{\text{sym}}^k \mathbf{X} \prod ((1 - \alpha)\tilde{\mathbf{A}}_{\text{sym}}^K + \alpha\mathbf{A}_{\text{sym}}^{K-1}) \mathbf{X}$
H ₂ GCN*	Coupled	$\prod_{k=0}^{2K} \mathbf{A}_{\text{sym}}^k \mathbf{X}$
GAT**	Attention	$\prod_{k=1}^K [\text{diag}(\mathbf{X}\mathbf{w}_{k,1})\tilde{\mathbf{A}} + \tilde{\mathbf{A}}\text{diag}(\mathbf{X}\mathbf{w}_{k,2})] \mathbf{X}$
DA-GNN**	Attention	$\sum_{k=0}^K \text{diag}(\tilde{\mathbf{A}}_{\text{sym}}^k \mathbf{X}\mathbf{w})\tilde{\mathbf{A}}_{\text{sym}}^k \mathbf{X}$

Q1: Reasons – Deep Dive



- Successes

✓ S1: Homophily $AX; A^2X; \dots \rightarrow$ neighbors' features

✗ ✓ S2: [Heterophily] AX : HURTS

A^2X : if concat., good

- Pain Points

✗ PP1: Lack of Robustness

$(\dots)X$



✗ PP2: Vulnerability to Noisy Features

$(\dots + I)X$



✗ PP3: Efficiency and Effectiveness 'too many cooks'

✗ PP4: Many Hyperparameters



'6 degrees'; $A^{100} \rightarrow$ rank-1



GNNEXP

Model	Type	Propagator function $\mathcal{P}(\mathbf{A}, \mathbf{X})$
LR	Linear	\mathbf{X}
SGC	Linear	$\tilde{\mathbf{A}}_{\text{sym}}^K \mathbf{X}$
DGC	Linear	$[(1 - T/K)\mathbf{I} + (T/K)\tilde{\mathbf{A}}_{\text{sym}}] \mathbf{X}$
$S^2\text{GC}$	Linear	$\sum_{k=1}^K (\alpha \mathbf{I} + (1 - \alpha)\tilde{\mathbf{A}}_{\text{sym}}^k) \mathbf{X}$
$G^2\text{CN}$	Linear	$\prod_{i=1}^N [\mathbf{I} - (T_i/K)(b_i - 1)\mathbf{I} + \mathbf{A}_{\text{sym}}] \mathbf{X}$
PPNP*	Decoupled	$(\mathbf{I} - (1 - \alpha)\tilde{\mathbf{A}}_{\text{sym}})^{-1} \mathbf{X}$
APPNP*	Decoupled	$[\sum_{k=0}^{K-1} \alpha(1 - \alpha)^k \tilde{\mathbf{A}}_{\text{sym}}^k + (1 - \alpha)^K \tilde{\mathbf{A}}_{\text{sym}}^K] \mathbf{X}$
GDC*	Decoupled	$\mathbf{S} = \text{sparse}_{\epsilon}(\sum_{k=0}^{\infty} (1 - \alpha)^k \tilde{\mathbf{A}}_{\text{sym}}^k)$ for $\tilde{\mathbf{S}}_{\text{sym}} \mathbf{X}$
GPR-GNN*	Decoupled	$\prod_{k=0}^K \tilde{\mathbf{A}}_{\text{sym}}^k \mathbf{X}$
ChebNet*	Coupled	$\prod_{k=0}^{K-1} \mathbf{A}_{\text{sym}}^k \mathbf{X}$
GCN*	Coupled	$\tilde{\mathbf{A}}_{\text{sym}}^K \mathbf{X}$
SAGE*	Coupled	$\prod_{k=0}^K \mathbf{A}_{\text{row}}^k \mathbf{X}$
GCNII*	Coupled	$\prod_{k=0}^{K-2} \tilde{\mathbf{A}}_{\text{sym}}^k \mathbf{X} \parallel ((1 - \alpha)\tilde{\mathbf{A}}_{\text{sym}}^K + \alpha\tilde{\mathbf{A}}_{\text{sym}}^{K-1}) \mathbf{X}$
$H_2\text{GCN}^*$	Coupled	$\prod_{k=0}^{2K} \mathbf{A}_{\text{sym}}^k \mathbf{X}$
GAT**	Attention	$\prod_{k=1}^K [\text{diag}(\mathbf{X}\mathbf{w}_{k,1})\tilde{\mathbf{A}} + \tilde{\mathbf{A}}\text{diag}(\mathbf{X}\mathbf{w}_{k,2})] \mathbf{X}$
DA-GNN**	Attention	$\sum_{k=0}^K \text{diag}(\tilde{\mathbf{A}}_{\text{sym}}^k \mathbf{X}\mathbf{w}) \tilde{\mathbf{A}}_{\text{sym}}^k \mathbf{X}$

Outline



- Problem Definition
- Q1: Reasons
- ➔ • Q2: Method
- Q3: Sanity Checks
- Experiments
- Conclusions

Q2: Method – Desiderata



- Desiderata and Design Decisions
 - D0: Linear Classifier
 - D1: Concatenating Winning Components
 - D2: Structure Features
 - D3: Orthogonalization and Sparsification
 - D4: Multi-Level Neighborhood Aggregation

Q2: Method – Desiderata

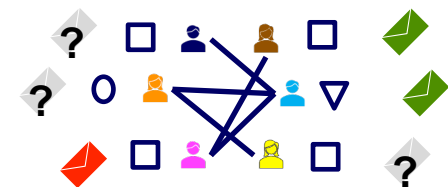
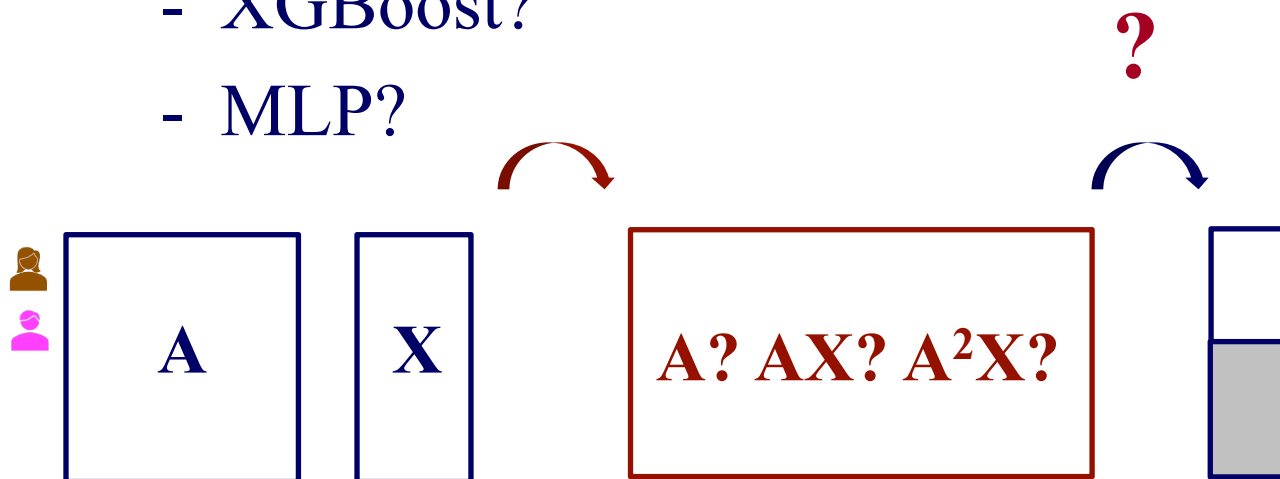


- Desiderata and Design Decisions
 - ➔ – D0: Linear Classifier
 - D1: Concatenating Winning Components
 - D2: Structure Features
 - D3: Orthogonalization and Sparsification
 - D4: Multi-Level Neighborhood Aggregation

Design Decision #0

Q: What classifier?

- Decision Trees?
- Random Forests?
- XGBoost?
- MLP?



Design Decision #0

A: Logistic regression (1-node MLP), after we find ‘good embeddings’

It’s simple and interpretable.

Logistic Regression (LR)



Design Decision #0

Q: # layers?

A: N/A

Q: Activation function?

A: Linear only (except the final LR step)



Q2: Method – Desiderata



- Desiderata and Design Decisions
 - D0: Linear Classifier
 - ➔ – D1: Concatenating Winning Components
 - ‘||’ not ‘+’
 - Heterophily
 - D2: Structure Features
 - D3: Orthogonalization and Sparsification
 - D4: Multi-Level Neighborhood Aggregation

Design Decision #1 – Concat.

- Concatenate \parallel ?
- Or add $+$?



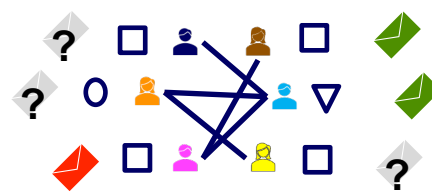
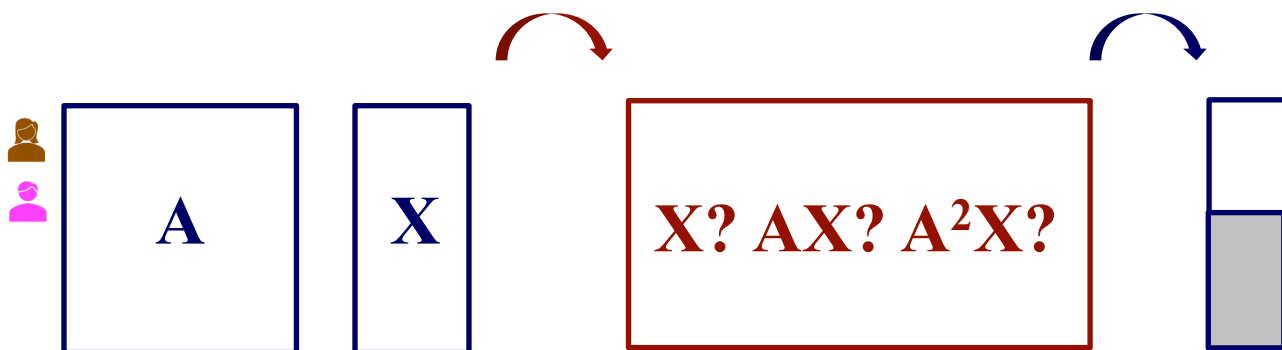
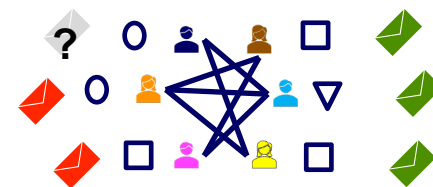
Design Decision #1 – Concat.

- Concatenate \parallel ? \rightarrow Heterophily / Robustness

$$AX \parallel A^2X \quad \checkmark$$

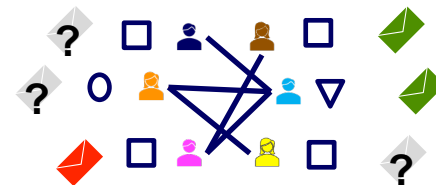
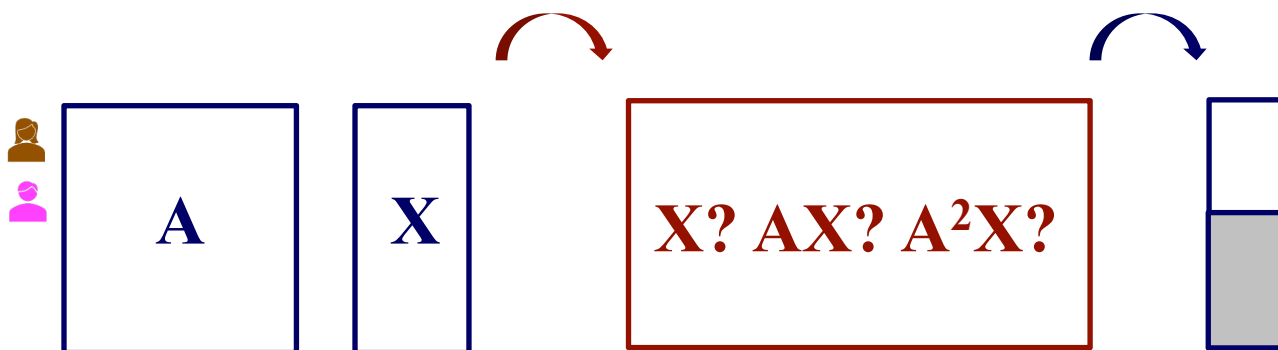
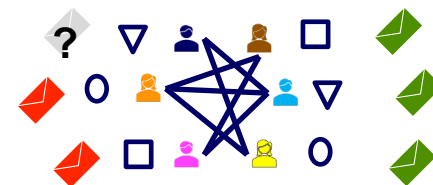
- Or add $+$?

$$AX + A^2X \quad \text{(with a red prohibition sign over it)}$$



Design Decision D1 – Hom. / Het.

- How to handle both homophily, AND heterophily?

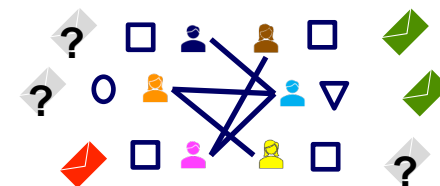
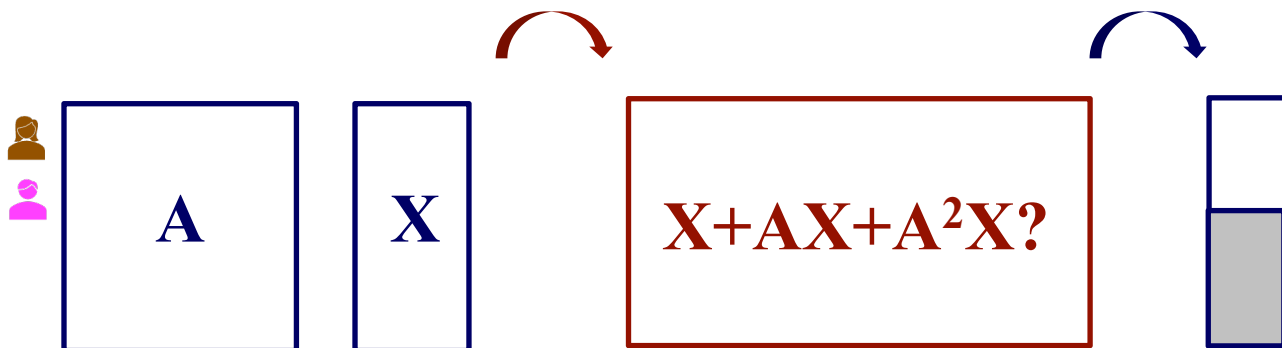
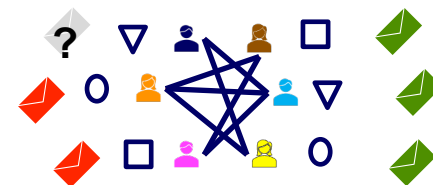


Design Decision D1 – Hom. / Het.



- How to handle both homophily, AND heterophily?

(If we add node features +
 1-, 2-, ... k-step neighbor features
 → mess and is hard to interpret)



Design Decision D1 – Hom. / Het.



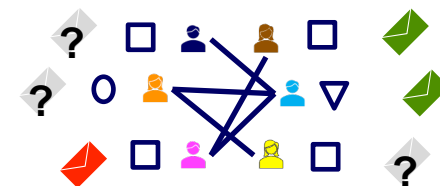
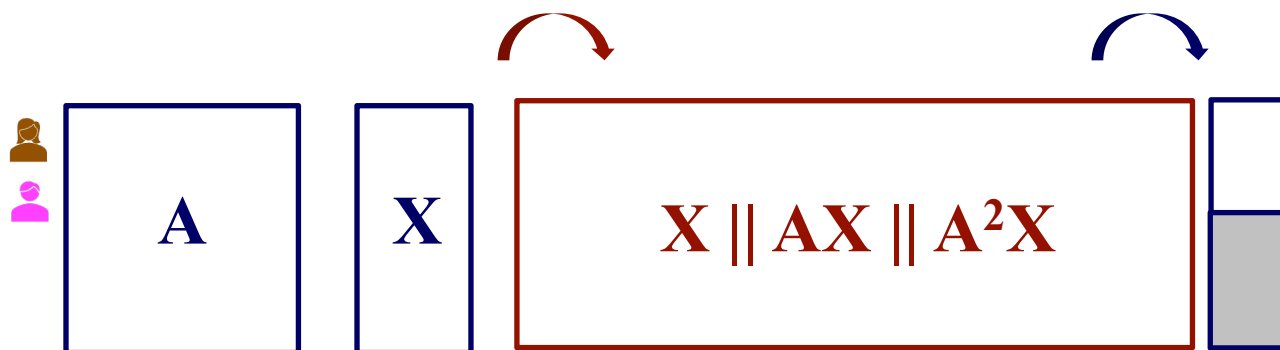
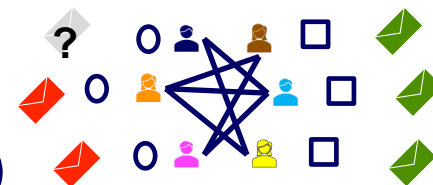
• How to handle both homophily, AND heterophily?



• Concatenate best homophily matrix,



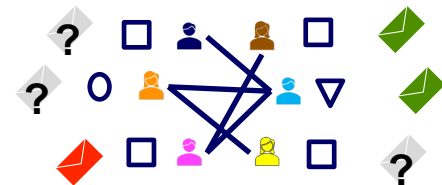
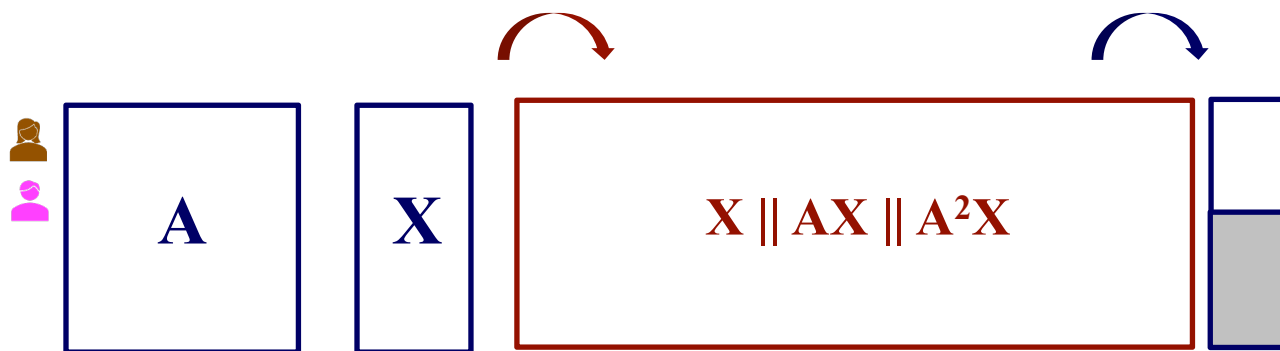
• AND grand-neighbors (even powers)



Design Decision D1 – Hom. / Het.



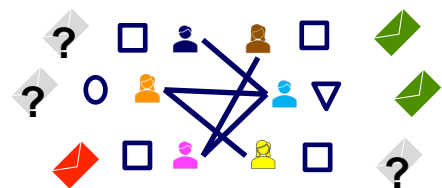
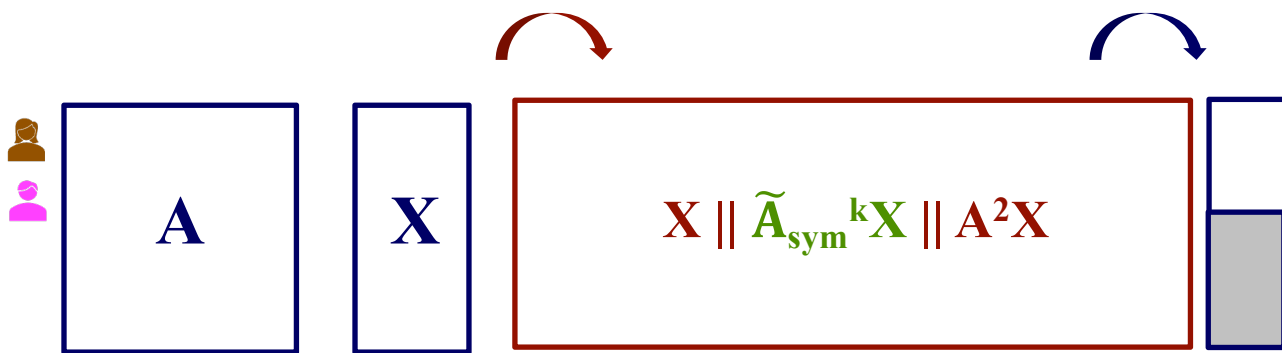
- Is A the best, for homophily?
- Is A^2 , heterophily?



Design Decision D1 – Hom. / Het.



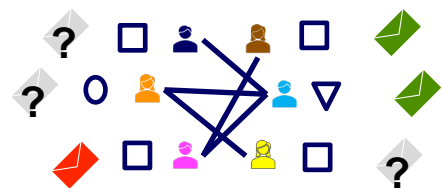
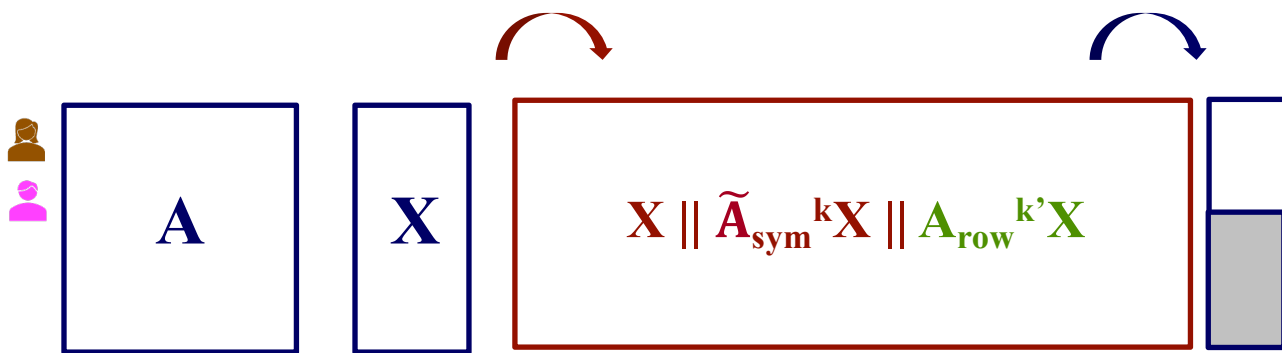
- Is A the best, for homophily?
- \tilde{A}_{sym} (symmetric norm. with self loops)
- Is A^2 , heterophily?



Design Decision D1 – Hom. / Het.



- Is A the best, for homophily?
- \tilde{A}_{sym} (symmetric norm. with self loops)
- Is A^2 , heterophily?
- A_{row} (row-norm; NO self loops)



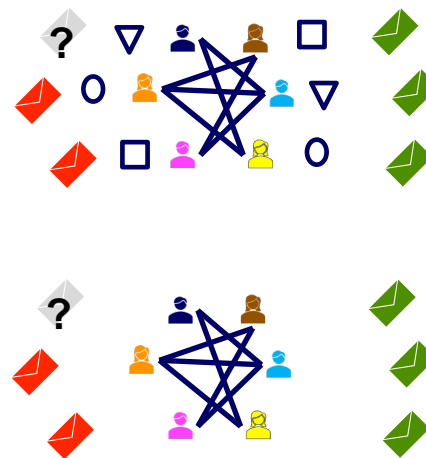
Q2: Method – Desiderata



- Desiderata and Design Decisions
 - D0: Linear Classifier
 - D1: Concatenating Winning Components
 - ➔ – D2: Structure Features
 - D3: Orthogonalization and Sparsification
 - D4: Multi-Level Neighborhood Aggregation

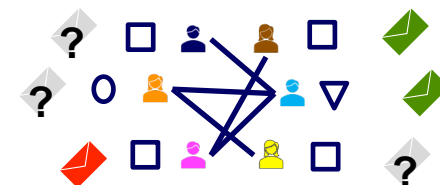
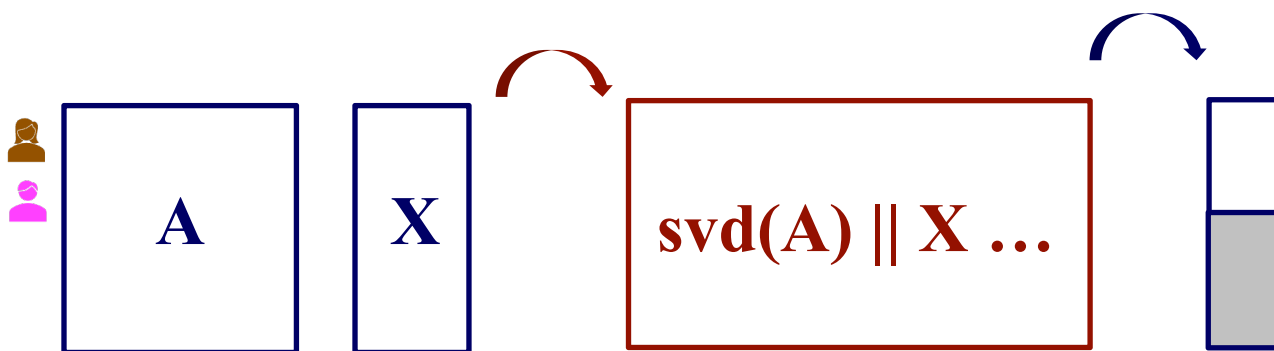
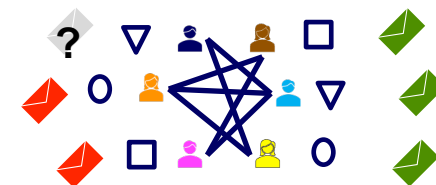
Design Decision D2 – Structure

- Q: What if (raw) features are useless?
 - i.e., uncorrelated to labels
 - E.g., hair color vs. voting pattern
 - Or, feature-less?



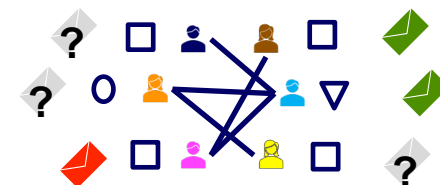
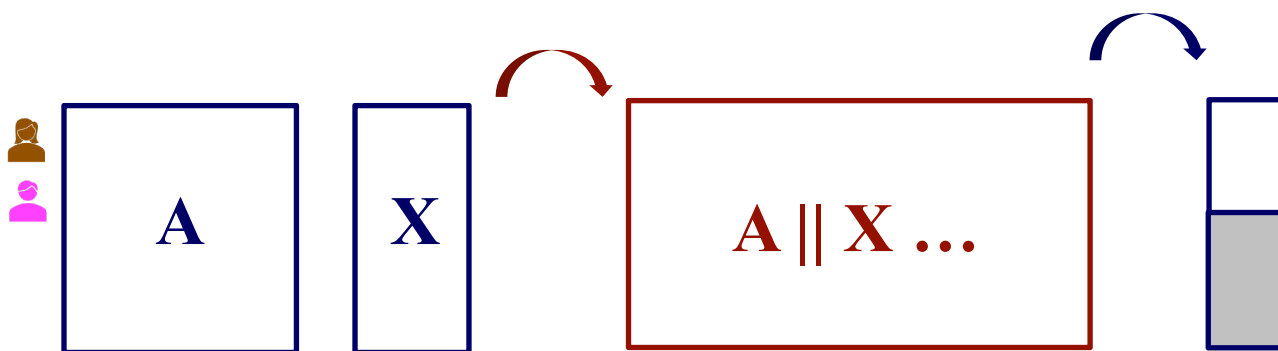
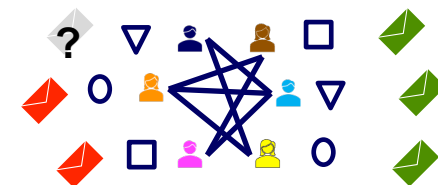
Design Decision D2 – Structure

- Q: What if (raw) features are useless?
- A: Add structure info., A



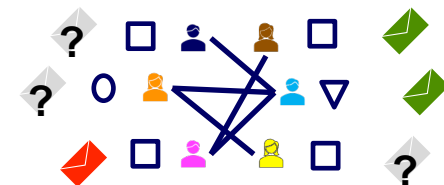
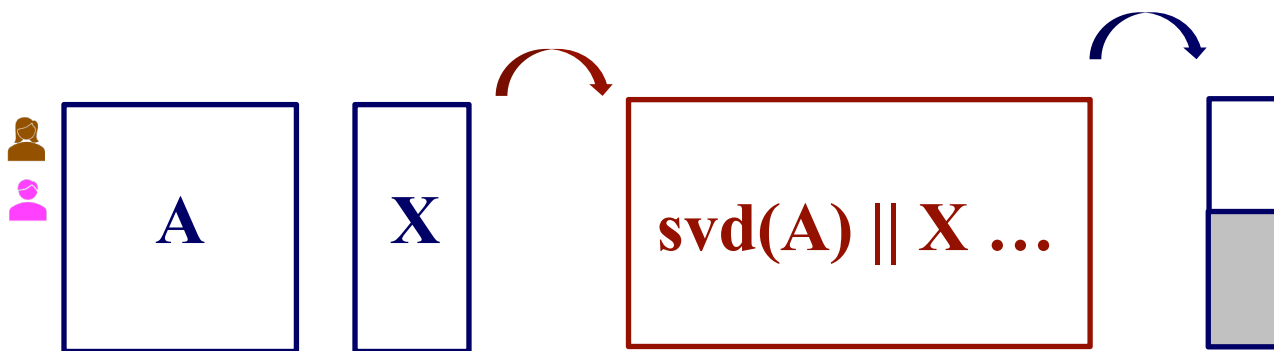
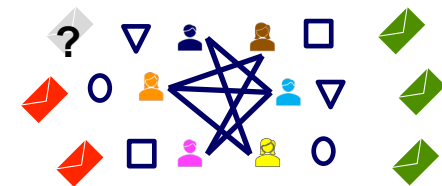
Design Decision D2 – Structure

- Q: What if (raw) features are useless?
- A: Add structure info., A
- (Q: but A is too wide, $1M \times 1M$)

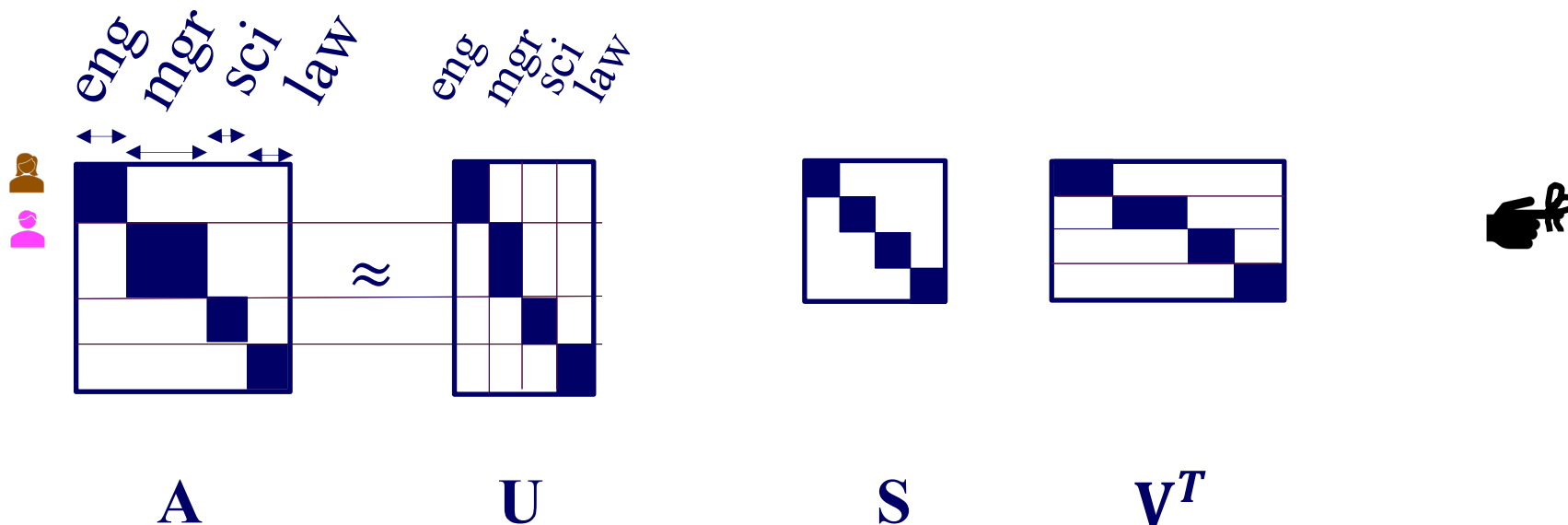


Design Decision D2 – Structure

- Q: What if (raw) features are useless?
- A: Add structure info., A
- (Q: but A is too wide, $1M \times 1M$)
- (A: U , where $U\Sigma V^T \approx A$ by SVD)

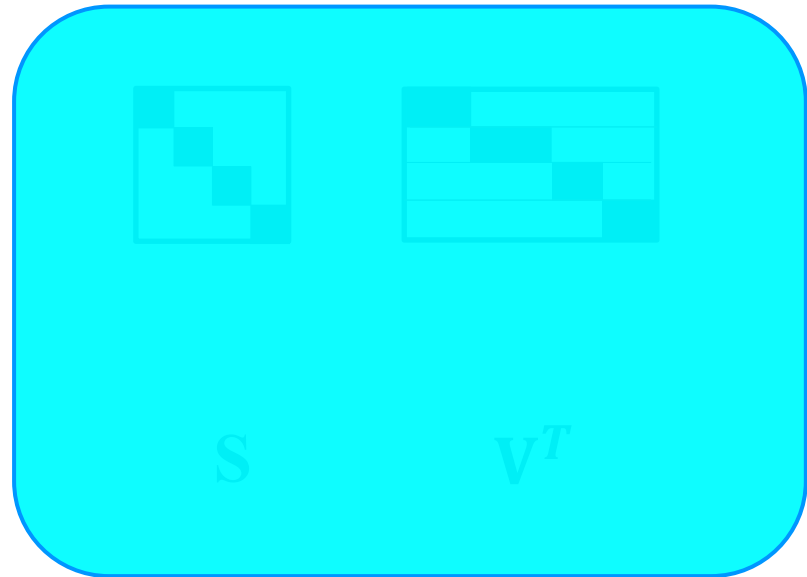
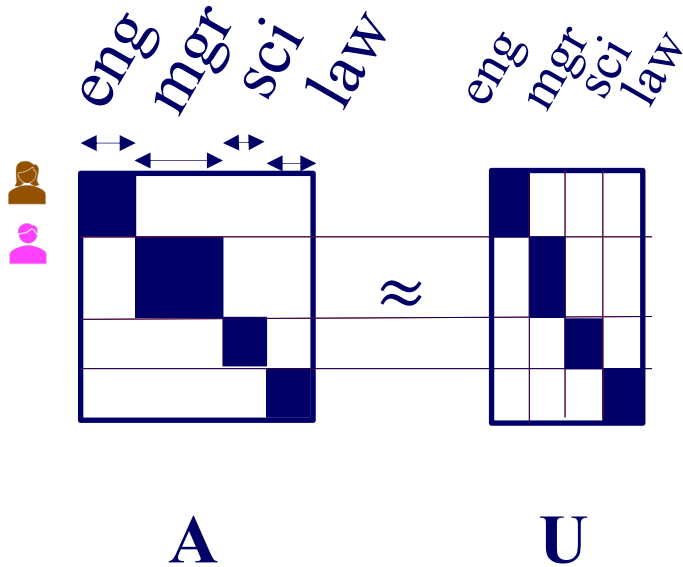


Reminder: SVD finds communities



Eg., if
block-diagonal =
communities ~
homophily

Reminder: SVD finds communities



Eg., if
block-diagonal =
communities ~
homophily

**Person x
community**

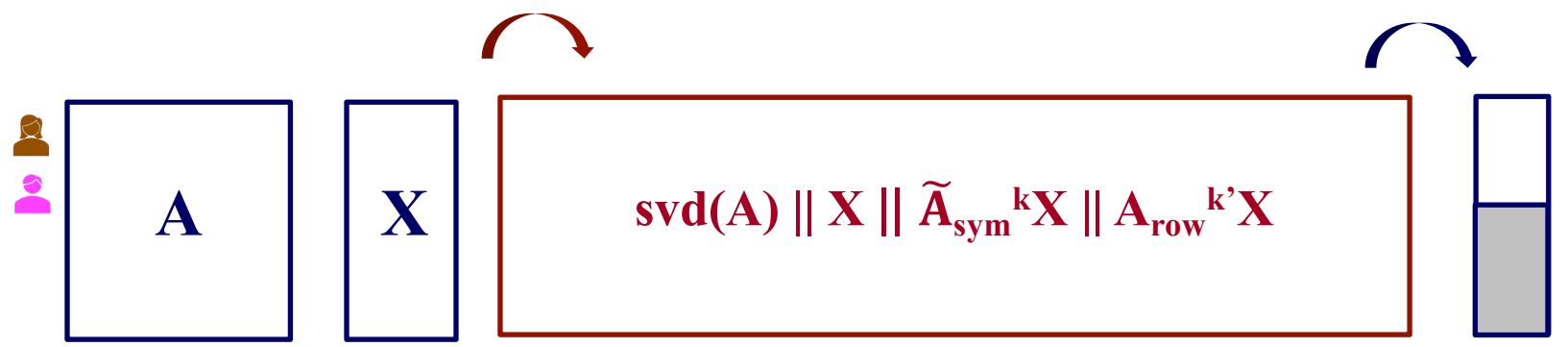
Q2: Method – Desiderata



- Desiderata and Design Decisions
 - D0: Linear Classifier
 - D1: Concatenating Winning Components
 - D2: Structure Features
 - ➔ – D3: Orthogonalization and Sparsification
 - Improve consistency
 - Remove redundant dimension(s)/component(s)
 - D4: Multi-Level Neighborhood Aggregation

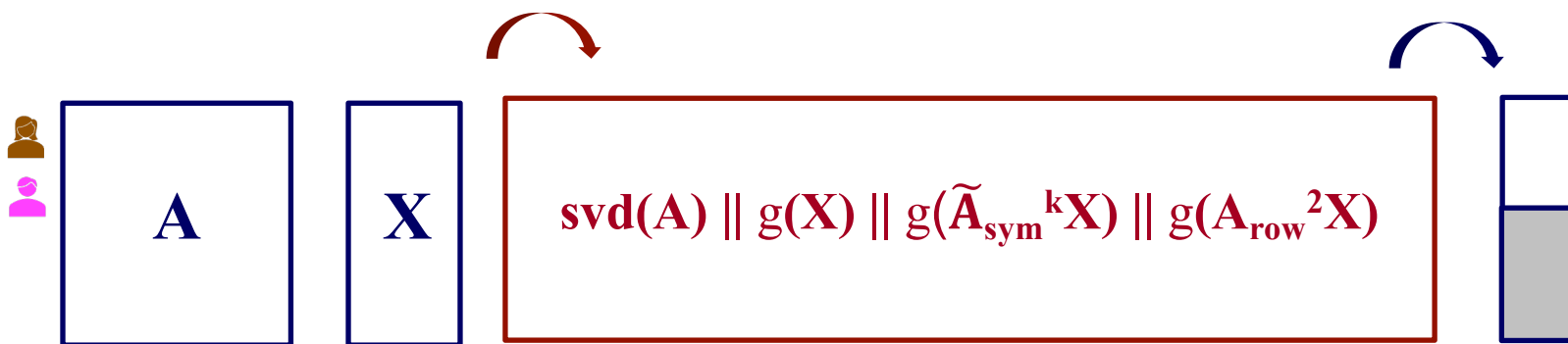
Design Decision D3 – Redundancy

- Q: Do we need all these ‘derived features’?



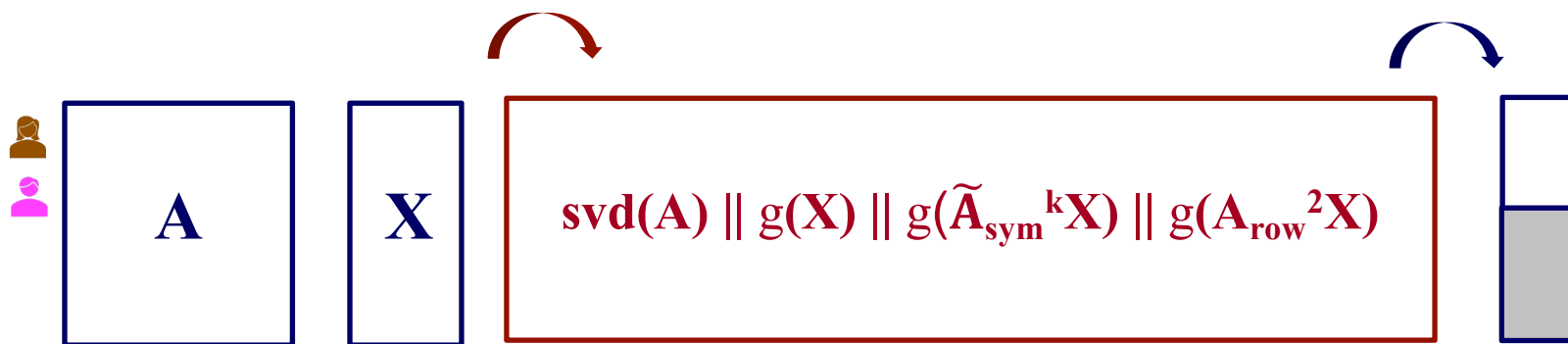
Design Decision D3 – Redundancy

- Q: Do we need all these ‘derived features’?
- A: No, dim. reduction $g()$ \rightarrow SVD/PCA on each component
 - Reduce correlated features
 - Fewer, good features \rightarrow more accurate



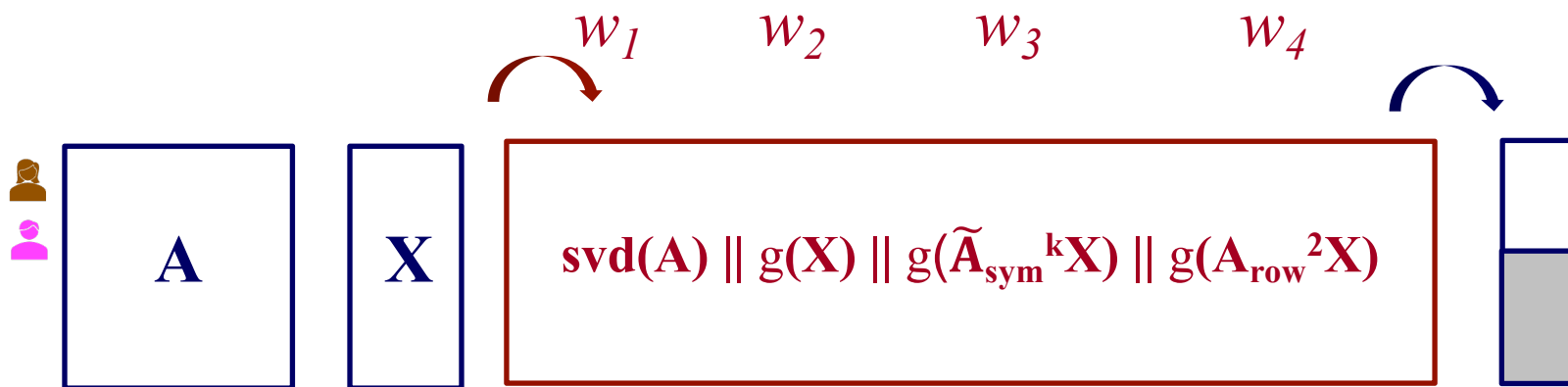
Design Decision D3 – More?

- Q: Do we always need all the components?



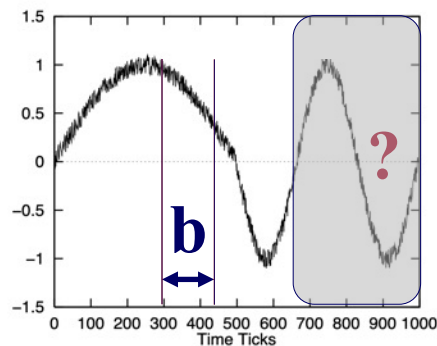
Design Decision D3 – More?

- Q: Do we always need all the components?
- A: Not necessarily, an entire component can be useless as well sometime



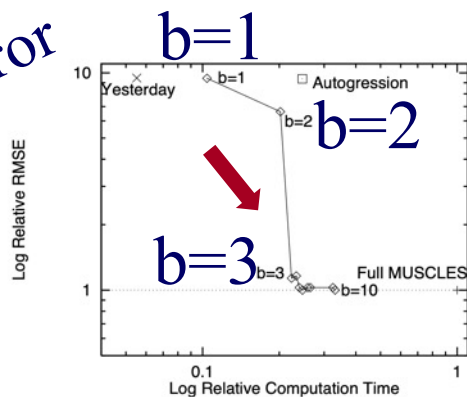


'Muscles' ICDE 2000



Common sense: *'the longer the window, the lower the error'*

error

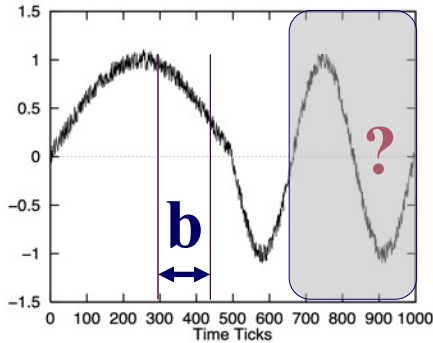


(a) US Dollar (CURRENCY)





'Muscles' ICDE 2000



Common sense: ~~the longer the window, the lower the error~~

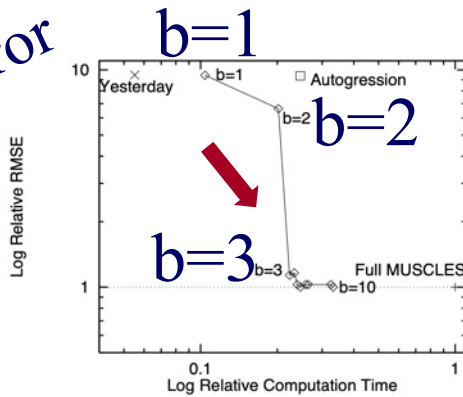


NOT always:

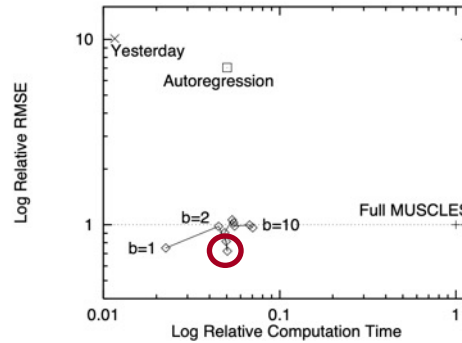
'too many cooks spoil the broth'



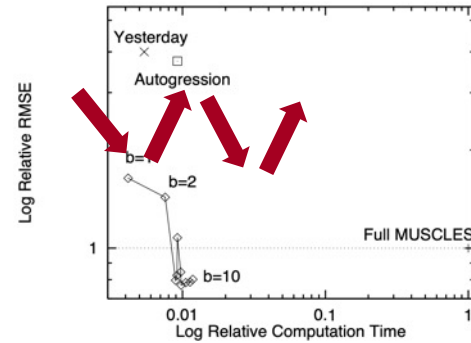
error



(a) US Dollar (CURRENCY)



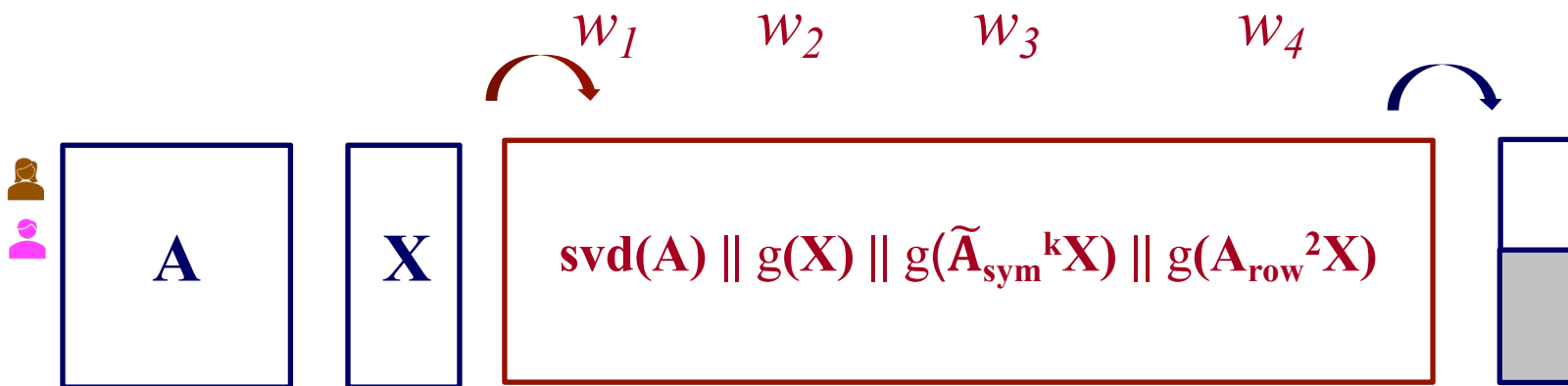
(b) 10-th modem (MODEM)



(c) 10-th stream (INTERNET)

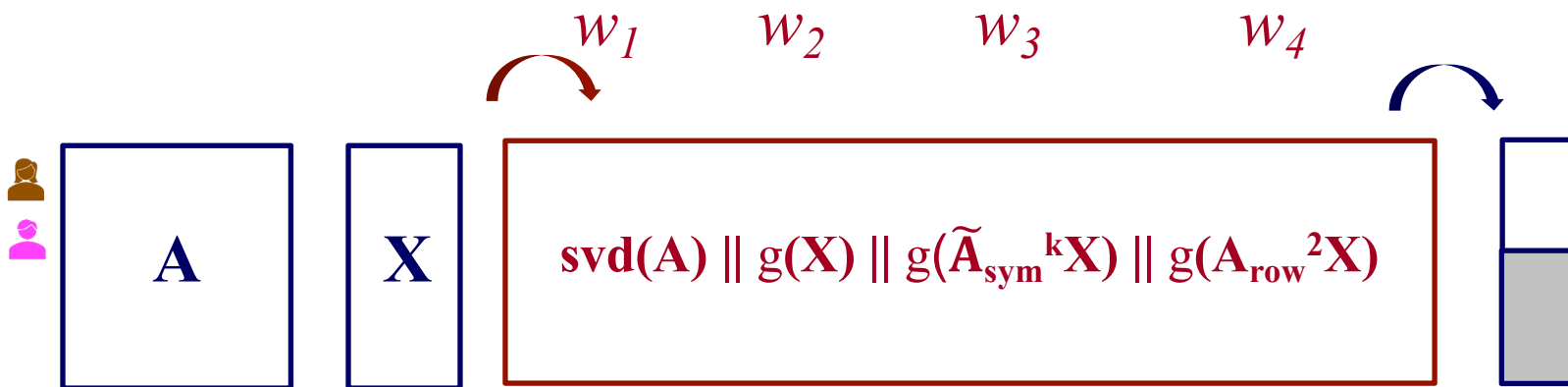
Design Decision D3 – More?

- Q: Do we always need all the components?
- A: Not necessarily, an entire component can be useless as well sometime
- A': Thus, 'group LASSO'



Design Decision D3 – More?

- Q: Do we always need all the components?
- A: Not necessarily, an entire component can be useless as well sometime
- A’: Thus, ‘group LASSO’
– Interpretable



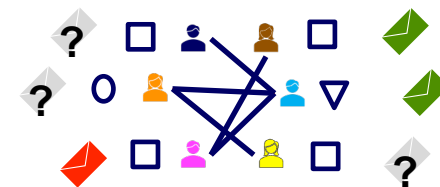
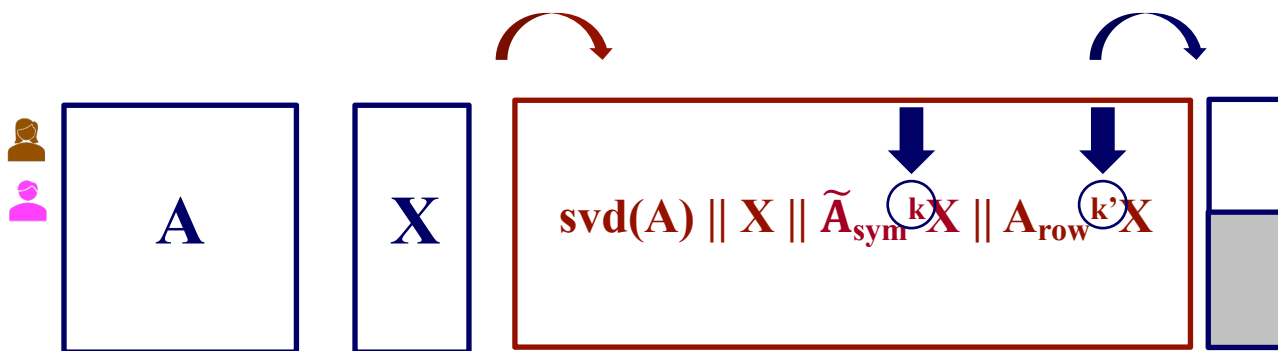
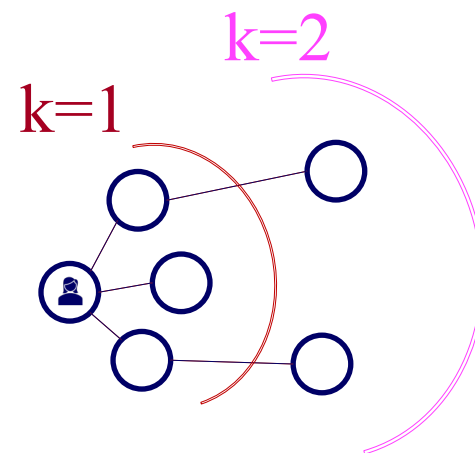
Q2: Method - Desiderata



- Desiderata and Design Decisions
 - D0: Linear Classifier
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 - D3: Orthogonalization and Sparsification
 - ➔ – D4: Multi-Level Neighborhood Aggregation

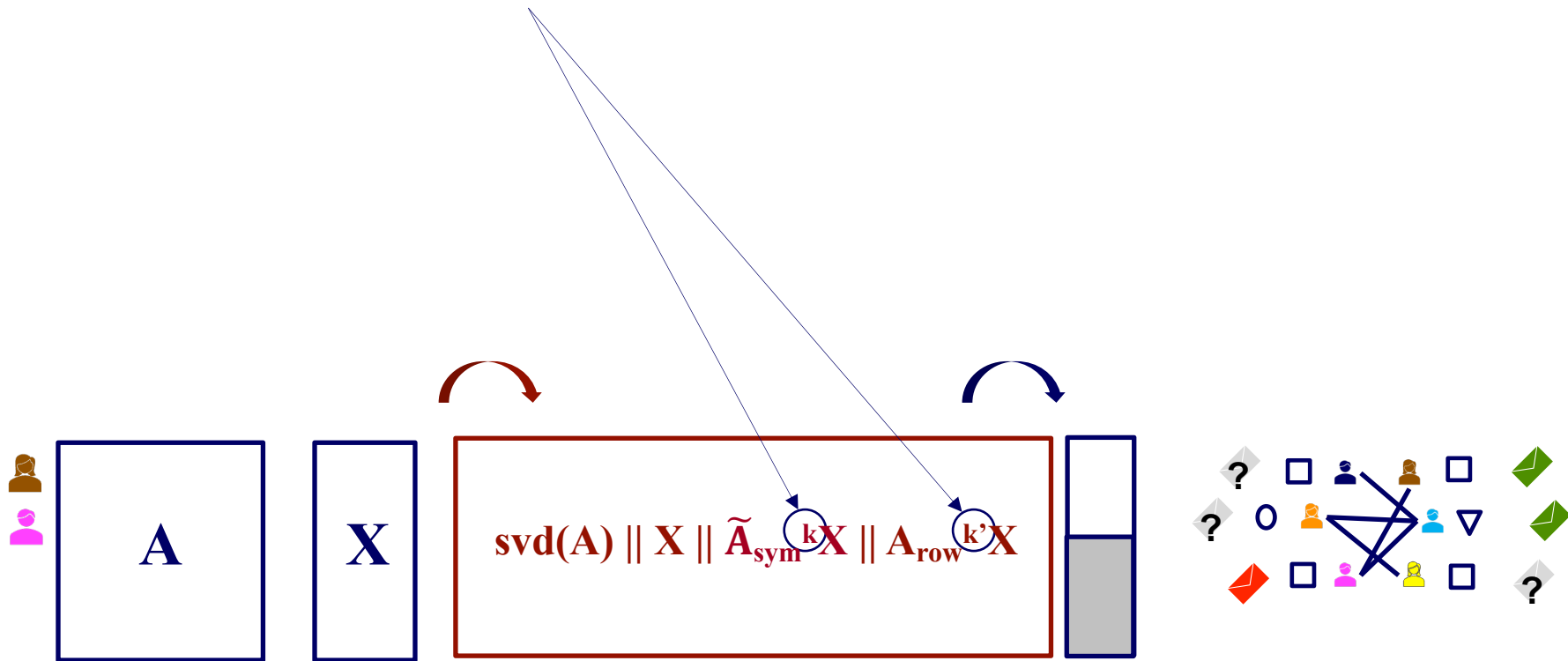
Design Decision D4 – Exponent

- Q: Influence horizon $k = ??$





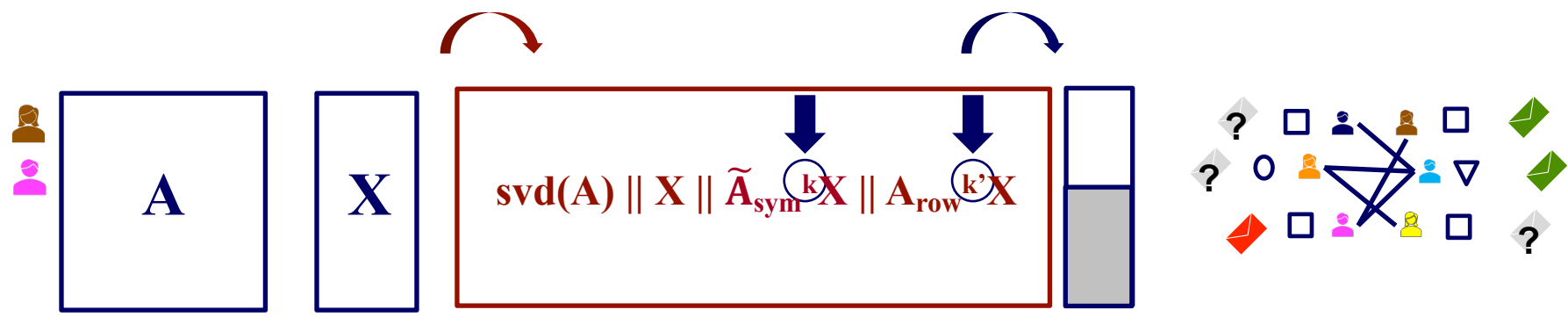
Design Decision D4 – Exponent

- Q: Influence horizon $k = ??$
- A: Propose: $k = 2$ (up to grand-neighbors)



Design Decision D4 – Exponent

- Q: Influence horizon $k = ??$
- A: Propose: $k = 2$ (up to grand-neighbors) 
- 0-, 1-, and 2-hop neighborhood
- b/c influence dissipates with distance
- b/c no over-smoothing 



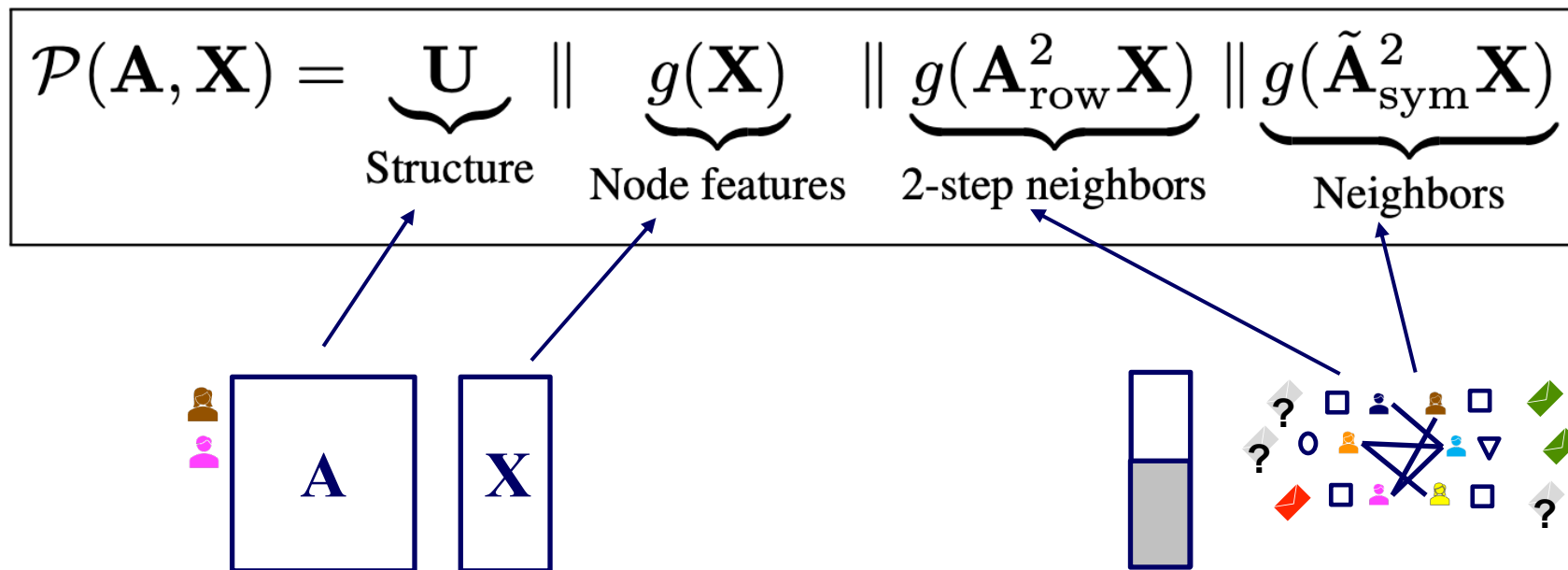
Q2: Method – SLIMG

4 components, carefully designed

$$\mathcal{P}(\mathbf{A}, \mathbf{X}) = \underbrace{\mathbf{U}}_{\text{Structure}} \parallel \underbrace{g(\mathbf{X})}_{\text{Node features}} \parallel \underbrace{g(\mathbf{A}_{\text{row}}^2 \mathbf{X})}_{\text{2-step neighbors}} \parallel \underbrace{g(\tilde{\mathbf{A}}_{\text{sym}}^2 \mathbf{X})}_{\text{Neighbors}}$$

Q2: Method – SLIMG

4 components, carefully designed



Q2: Method – SLIMG

4 components, carefully designed

$$\mathcal{P}(\mathbf{A}, \mathbf{X}) = \underbrace{\mathbf{U}}_{\text{Structure}} \parallel \underbrace{g(\mathbf{X})}_{\text{Node features}} \parallel \underbrace{g(\mathbf{A}_{\text{row}}^2 \mathbf{X})}_{\text{2-step neighbors}} \parallel \underbrace{g(\tilde{\mathbf{A}}_{\text{sym}}^2 \mathbf{X})}_{\text{Neighbors}}$$

Orthogonalization:
 PCA

Q2: Method – SLIMG

“Careful Simplicity”



4 components, carefully designed

$$\mathcal{P}(\mathbf{A}, \mathbf{X}) = \underbrace{\mathbf{U}}_{\text{Structure}} \parallel \underbrace{g(\mathbf{X})}_{\text{Node features}} \parallel \underbrace{g(\mathbf{A}_{\text{row}}^2 \mathbf{X})}_{\text{2-step neighbors}} \parallel \underbrace{g(\tilde{\mathbf{A}}_{\text{sym}}^2 \mathbf{X})}_{\text{Neighbors}}$$

Outline



- Problem Definition
- Q1: Reasons
- Q2: Method
- ➔ • Q3: Sanity Checks
- Experiments
- Conclusions

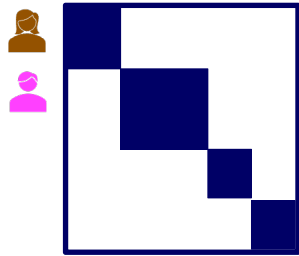
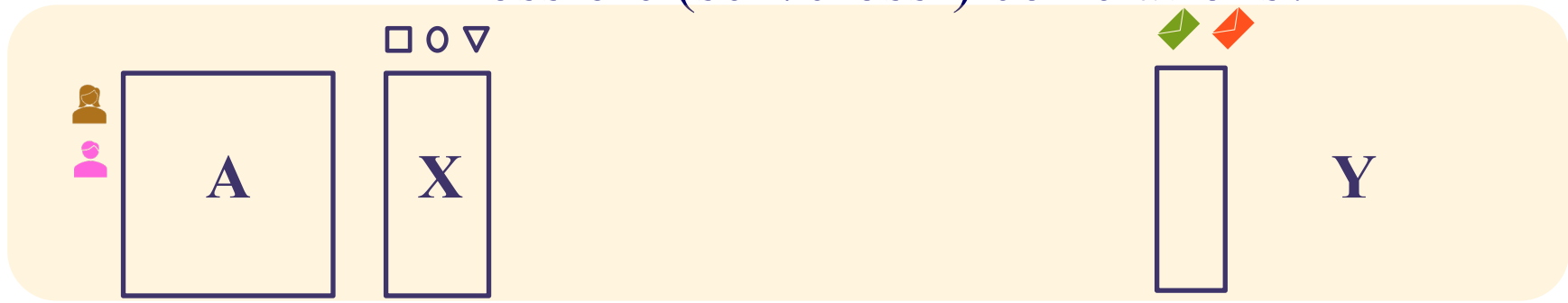
Q3: Sanity Checks

- Q: What are possible settings?
- A: Cross-product of scenarios

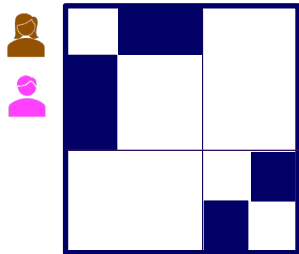
Q3: Sanity Checks

- Q: What are possible settings?
- A: Cross-product of scenarios
 - S1: connectivity
 - Block-diagonal
 - Block-off-diagonal
 - Random

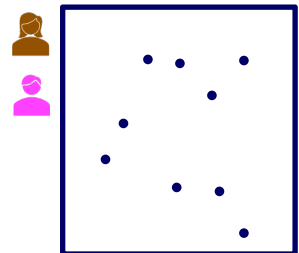
Possible (self/cross-) correlations?



block-diagonal =
communities ~
homophily



block-off-diagonal =
'familiar strangers' ~
heterophily

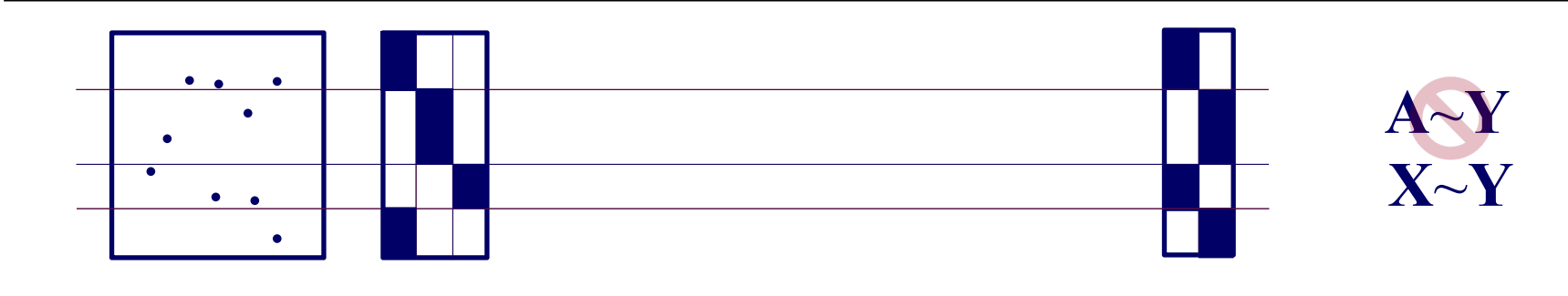
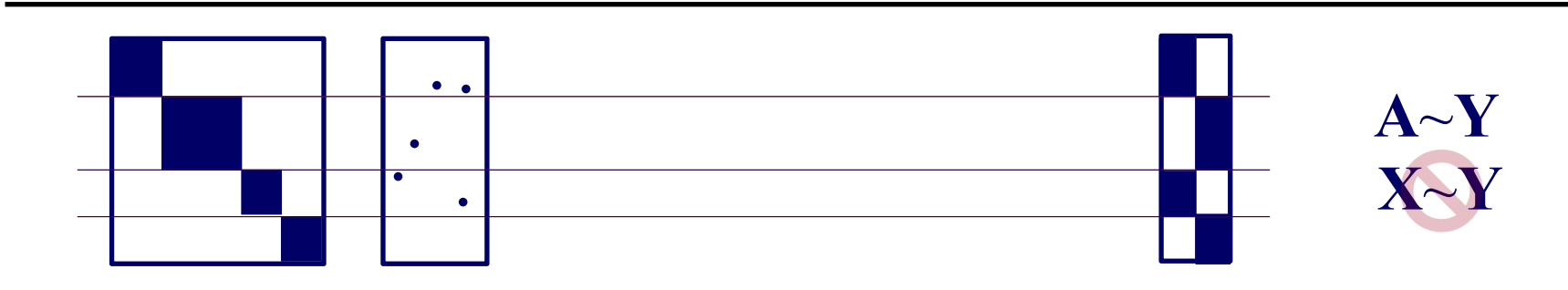
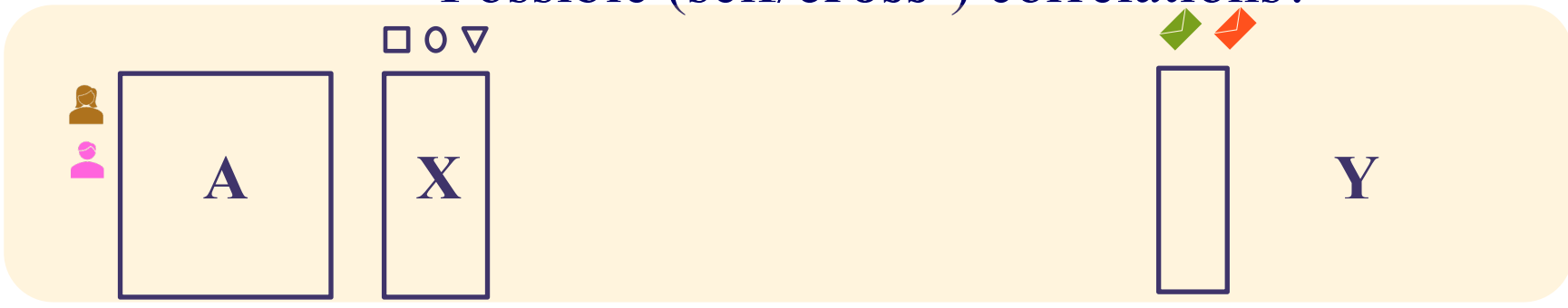


Random

Q3: Sanity Checks

- Q: What are possible settings?
- A: Cross-product of scenarios
 - S1: connectivity
 - S2: cross-correlations – labels are correlated w/
 - Connectivity only (**A**)
 - Features only (**X**)
 - Both
 - (None)

Possible (self/cross-) correlations?



(omitted – obvious – everybody does well)

$A \sim Y$
 $X \sim Y$

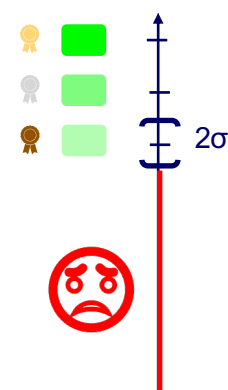
Q3: Sanity Checks

Only X
helps

Only A
helps

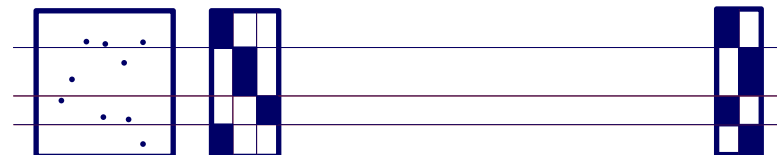
Both A, X
help

Model	Only X helps	Only A helps		Both X and A help				Avg. Acc	Avg. Rank
	Semantic X Uniform A	Random X Homophily	Random X Heterophily	Structural X Homophily	Structural X Heterophily	Semantic X Homophily	Semantic X Heterophily		
LR	83.7±0.6	24.2±0.7	24.2±0.7	71.4±0.9	66.8±2.2	83.4±0.6	83.4±0.6	62.4 (26.9)	10.7 (5.4)
Reg. Kernel	82.7±0.5	27.9±0.4	24.3±1.0	75.7±0.2	65.3±1.6	91.5±0.5	79.5±0.3	63.8 (27.0)	10.4 (4.3)
Diff. Kernel	26.8±1.7	38.0±8.7	37.6±7.5	79.5±0.3	73.5±0.6	70.9±23.	56.1±27.	54.6 (20.7)	10.6 (4.0)
RW Kernel	72.2±0.7	37.0±0.4	24.5±1.3	81.3±1.2	51.0±1.1	94.5±0.9	57.8±0.7	59.8 (24.7)	10.4 (3.6)
SGC	44.6±9.8	64.3±0.7	50.2±14.	87.1±0.6	84.3±0.5	93.9±0.9	91.5±0.5	73.7 (20.4)	5.7 (3.1)
DGC	63.8±1.0	50.5±13.	26.0±0.9	88.6±1.0	45.3±1.3	96.2±0.4	54.0±0.6	60.6 (24.6)	8.3 (5.9)
S ² GC	79.9±0.6	38.5±12.	25.4±0.9	88.4±1.0	67.9±1.5	95.9±0.6	78.0±0.5	67.7 (26.2)	7.4 (3.4)
G ² CN	25.2±0.3	24.2±1.1	25.0±0.1	88.5±1.0	88.6±1.2	24.3±1.1	50.7±31.	46.6 (30.2)	11.6 (6.3)
GCN	36.3±3.5	46.7±8.0	43.7±1.9	83.3±1.3	72.2±1.7	91.2±1.2	80.3±3.9	64.8 (22.1)	8.1 (3.0)
SAGE	80.3±1.1	31.1±0.7	34.6±2.1	83.9±0.8	81.3±0.7	94.4±0.5	94.4±0.9	71.4 (27.0)	5.7 (2.9)
GCNII	73.5±1.2	30.7±0.7	27.1±1.3	84.2±0.8	69.0±1.4	90.6±0.9	80.4±1.2	65.1 (25.7)	8.7 (1.8)
H ² GCN	80.2±1.5	27.0±1.0	27.5±0.8	78.0±0.9	74.6±1.3	91.9±0.7	92.2±0.9	67.3 (28.2)	8.0 (3.9)
APPNP	66.0±2.6	30.3±1.2	25.2±0.7	71.2±4.9	43.8±2.0	83.2±3.8	58.7±4.5	54.1 (21.6)	12.9 (2.0)
GPR-GNN	73.4±0.4	74.6±0.7	65.9±2.1	89.9±0.6	87.6±1.2	95.0±1.1	91.9±1.1	82.6 (11.2)	3.3 (2.1)
GAT	32.7±5.5	42.6±4.8	36.8±5.7	64.0±5.7	55.6±6.8	68.5±7.1	67.0±12.	52.5 (15.0)	11.6 (4.1)
SLIMG (Ours)	81.0±1.1	87.1±1.4	89.2±1.2	88.1±0.5	88.9±0.7	94.4±0.6	93.9±0.5	88.9 (4.5)	2.6 (1.8)



Q3: Sanity Checks – Part 1/3

Only X helps

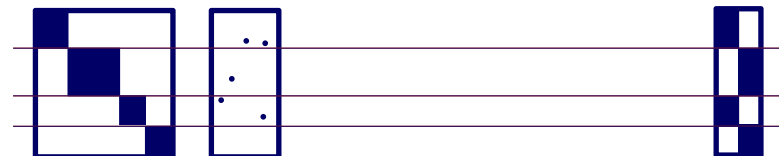


Model	Only X helps Semantic X Uniform A
LR	83.7±0.6
Reg. Kernel	82.7±0.5
Diff. Kernel	26.8±1.7
RW Kernel	72.2±0.7
SGC	44.6±9.8
DGC	63.8±1.0
S ² GC	79.9±0.6
G ² CN	25.2±0.3
GCN	36.3±3.5
SAGE	80.3±1.1
GCNII	73.5±1.2
H ² GCN	80.2±1.5
APPNP	66.0±2.6
GPR-GNN	73.4±0.4
GAT	32.7±5.5
SLIMG (Ours)	81.0±1.1

- 1) Logistic Reg. does great
- 2) ... as well as SLIMG
- 3) Rest: confused
 - 1) ... often 1/4 (random)

Q3: Sanity Checks – Part 2/3

Only A helps



Model	Only A helps	
	Random X Homophily	Random X Heterophily
LR	24.2±0.7	24.2±0.7
Reg. Kernel	27.9±0.4	24.3±1.0
Diff. Kernel	38.0±8.7	37.6±7.5
RW Kernel	37.0±0.4	24.5±1.3
SGC	64.3±0.7	50.2±14.
DGC	50.5±13.	26.0±0.9
S ² GC	38.5±12.	25.4±0.9
G ² CN	24.2±1.1	25.0±0.1
GCN	46.7±8.0	43.7±1.9
SAGE	31.1±0.7	34.6±2.1
GCNII	30.7±0.7	27.1±1.3
H ² GCN	27.0±1.0	27.5±0.8
APPNP	30.3±1.2	25.2±0.7
GPR-GNN	74.6±0.7	65.9±2.1
GAT	42.6±4.8	36.8±5.7
SLIMG (Ours)	87.1±1.4	89.2±1.2

As expected:

- 1) **Only** SLIMG works
- 2) All else fail

$$(A^k + A^j + \dots)X \parallel (A^{k'} \dots)X$$

Q3: Sanity Checks – Part 3/3

Both A, X
help

Model	Both X and A help			
	Structural X Homophily	Structural X Heterophily	Semantic X Homophily	Semantic X Heterophily
LR	71.4±0.9	66.8±2.2	83.4±0.6	83.4±0.6
Reg. Kernel	75.7±0.2	65.3±1.6	91.5±0.5	79.5±0.3
Diff. Kernel	79.5±0.3	73.5±0.6	70.9±23.	56.1±27.
RW Kernel	81.3±1.2	51.0±1.1	94.5±0.9	57.8±0.7
SGC	87.1±0.6	84.3±0.5	93.9±0.9	91.5±0.5
DGC	88.6±1.0	45.3±1.3	96.2±0.4	54.0±0.6
S ² GC	88.4±1.0	67.9±1.5	95.9±0.6	78.0±0.5
G ² CN	88.5±1.0	88.6±1.2	24.3±1.1	50.7±31.
GCN	83.3±1.3	72.2±1.7	91.2±1.2	80.3±3.9
SAGE	83.9±0.8	81.3±0.7	94.4±0.5	94.4±0.9
GCNII	84.2±0.8	69.0±1.4	90.6±0.9	80.4±1.2
H ² GCN	78.0±0.9	74.6±1.3	91.9±0.7	92.2±0.9
APPNP	71.2±4.9	43.8±2.0	83.2±3.8	58.7±4.5
GPR-GNN	89.9±0.6	87.6±1.2	95.0±1.1	91.9±1.1
GAT	64.0±5.7	55.6±6.8	68.5±7.1	67.0±12.
SLIMG (Ours)	88.1±0.5	88.9±0.7	94.4±0.6	93.9±0.5

- 1) Several do OK on hom.
- 2) Fewer work on het.
- 3) SLIMG: never ‘red’

Outline



- Problem Definition
- Q1: Reasons
- Q2: Method
- Q3: Sanity Checks
- ➔ • Experiments
 - E1: Accurate & Robust
 - E2: Interpretable
 - E3: Fast & Scalable
- Conclusions

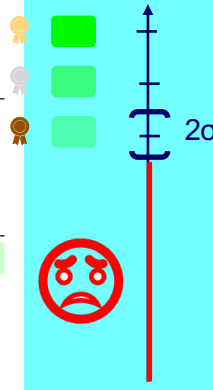
E1: Accurate & Robust Real-World Datasets



7 Homophily

6 Heterophily

Model	Cora	CiteSeer	PubMed	Comp.	Photo	ArXiv	Products	Cham.	Squirrel	Actor	Penn94	Twitch	Pokec	Avg. Rank
LR	51.5±1.2	52.9±4.5	79.9±0.5	73.9±1.2	79.3±1.5	48.3±1.9	56.4±0.5	24.9±1.7	26.7±1.9	27.8±0.8	63.5±0.5	53.0±0.1	61.3±0.0	11.7 (4.2)
Reg. Kernel	67.8±2.5	62.1±4.4	83.4±1.4	80.3±1.4	87.1±1.2	O.O.M.	O.O.M.	29.4±2.6	24.3±2.3	29.6±1.4	O.O.M.	O.O.M.	O.O.M.	12.2 (3.8)
Diff. Kernel	70.6±1.5	62.7±3.8	82.1±0.4	83.1±1.0	89.8±0.6	O.O.M.	O.O.M.	34.5±7.9	28.3±1.5	24.7±0.9	53.5±0.8	O.O.M.	O.O.M.	11.8 (2.5)
RW Kernel	72.7±1.7	64.1±3.9	83.1±0.7	84.2±0.7	90.6±0.7	63.2±0.2	74.2±0.0	34.9±3.5	25.0±1.6	26.4±1.1	63.1±0.7	57.6±0.1	59.5±0.0	8.3 (3.3)
SGC	76.2±1.1	65.8±3.9	84.1±0.8	83.7±1.6	90.1±0.9	65.0±3.4	74.6±5.1	38.1±4.5	33.1±1.0	24.6±0.8	64.0±1.1	56.5±0.1	69.8±0.0	6.6 (4.2)
DGC	77.8±1.4	66.1±4.2	84.3±0.6	83.9±0.7	90.4±0.2	65.2±4.0	68.7±13.	37.2±3.7	29.2±1.2	25.2±2.1	62.5±0.4	58.2±0.2	60.7±0.1	6.6 (3.2)
S ² GC	78.3±1.5	66.9±4.4	84.3±0.3	83.1±0.8	90.1±0.8	62.0±7.4	58.3±18.	34.9±4.9	27.6±1.8	26.7±1.8	63.1±0.5	58.7±0.1	61.2±0.0	6.6 (2.7)
G ² CN	76.6±1.5	64.2±3.3	81.4±0.6	82.8±1.6	88.8±0.5	O.O.M.	O.O.M.	40.7±2.9	32.1±1.5	24.3±0.5	O.O.M.	O.O.M.	O.O.M.	10.5 (4.5)
GCN	76.0±1.2	65.0±2.9	84.3±0.5	85.1±0.9	91.6±0.5	62.8±0.6	O.O.M.	38.5±3.0	31.4±1.8	26.8±0.4	62.9±0.7	57.0±0.1	63.9±0.4	6.3 (2.4)
SAGE	74.6±1.3	63.7±3.6	82.9±0.4	83.8±0.5	90.6±0.5	61.5±0.6	O.O.M.	39.8±4.3	27.0±1.3	27.8±0.9	O.O.M.	56.6±0.4	68.9±0.1	8.5 (3.5)
GCNII	77.8±1.7	63.4±3.0	84.9±0.8	82.3±1.8	90.8±0.6	45.7±0.5	O.O.M.	30.5±2.5	21.9±3.0	29.0±1.3	64.5±0.5	56.9±0.6	62.1±0.3	8.4 (4.6)
H ² GCN	77.6±0.9	64.7±3.8	85.4±0.4	49.5±16.	75.8±11.	O.O.M.	O.O.M.	31.9±2.6	25.0±0.5	28.9±0.6	63.9±0.4	58.7±0.0	O.O.M.	8.9 (4.9)
APPNP	80.0±0.6	67.1±2.8	84.6±0.5	84.2±1.7	92.5±0.3	53.4±1.3	O.O.M.	30.9±4.7	23.9±3.2	26.1±1.0	63.7±0.9	47.3±0.3	57.4±0.4	7.6 (4.8)
GPR-GNN	78.8±1.3	64.2±4.0	85.1±0.7	85.0±1.0	92.6±0.3	58.5±0.8	O.O.M.	31.7±4.7	26.2±1.6	29.5±1.1	64.5±0.4	57.6±0.2	67.6±0.1	5.4 (3.7)
GAT	78.2±1.2	65.8±4.0	83.6±0.2	85.4±1.4	91.7±0.5	58.2±1.0	O.O.M.	39.1±4.1	28.6±0.6	26.4±0.4	60.5±0.8	O.O.M.	O.O.M.	7.5 (3.7)
SLIMG	77.8±1.1	67.1±2.3	84.6±0.5	86.3±0.7	91.8±0.5	66.3±0.3	84.9±0.0	40.8±3.2	31.1±0.7	30.9±0.6	68.2±0.6	59.7±0.1	73.9±0.1	1.9 (1.5)



E1: Accurate & Robust Real-World Datasets

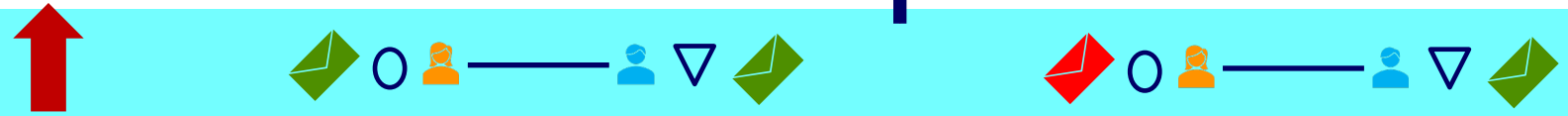
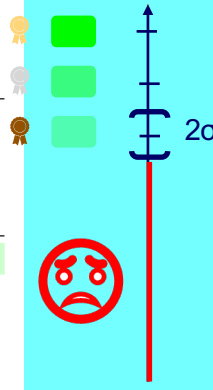


7 Homophily

6 Heterophily

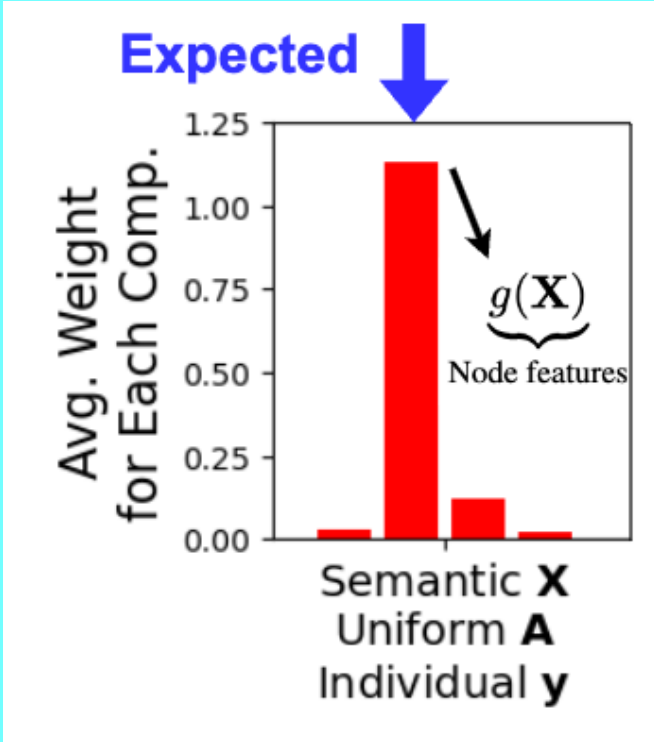
Many O.O.M.

Model	Cora	CiteSeer	PubM	ArXiv	Products	Cham.	Squirrel	Actor	Penn94	Twitch	Pokec	Avg. Rank		
LR	51.5±1.2	52.9±4.5	59.3±1.5	48.3±1.9	56.4±0.5	24.9±1.7	26.7±1.9	27.8±0.8	63.5±0.5	53.0±0.1	61.3±0.0	11.7 (4.2)		
Reg. Kernel	67.8±2.5	62.1±1.4	87.1±1.2	O.O.M.	O.O.M.	29.4±2.6	24.3±2.3	29.6±1.4	O.O.M.	O.O.M.	O.O.M.	12.2 (3.8)		
Diff. Kernel	70.6±1.5	83.1±1.0	89.8±0.6	O.O.M.	O.O.M.	34.5±7.9	28.3±1.5	24.7±0.9	53.5±0.8	O.O.M.	O.O.M.	11.8 (2.5)		
RW Kernel	72.1±0.7	84.2±0.7	90.6±0.7	63.2±0.2	74.2±0.0	34.9±3.5	25.0±1.6	26.4±1.1	63.1±0.7	57.6±0.1	59.5±0.0	8.3 (3.3)		
SGC			84.1±0.8	83.7±1.6	90.1±0.9	65.0±3.4	74.6±5.1	38.1±4.5	33.1±1.0	24.6±0.8	64.0±1.1	56.5±0.1	69.8±0.0	6.6 (4.2)
DCN		61.1±4.2	84.3±0.6	83.9±0.7	90.4±0.2	65.2±4.0	68.7±13.	37.2±3.7	29.2±1.2	25.2±2.1	62.5±0.4	58.2±0.2	60.7±0.1	6.6 (3.2)
		66.9±4.4	84.3±0.3	83.1±0.8	90.1±0.8	62.0±7.4	58.3±18.	34.9±4.9	27.6±1.8	26.7±1.8	63.1±0.5	58.7±0.1	61.2±0.0	6.6 (2.7)
	60±1.5	64.2±3.3	81.4±0.6	82.8±1.6	88.8±0.5	O.O.M.	O.O.M.	40.7±2.9	32.1±1.5	24.3±0.5	O.O.M.	O.O.M.	O.O.M.	10.5 (4.5)
GCN	76.0±1.2	65.0±2.9	84.3±0.5	85.1±0.9	91.6±0.5	62.8±0.6	O.O.M.	38.5±3.0	31.4±1.8	26.8±0.4	62.9±0.7	57.0±0.1	63.9±0.4	6.3 (2.4)
SAGE	74.6±1.3	63.7±3.6	82.9±0.4	83.8±0.5	90.6±0.5	61.5±0.6	O.O.M.	39.8±4.3	27.0±1.3	27.8±0.9	O.O.M.	56.6±0.4	68.9±0.1	8.5 (3.5)
GCNII	77.8±1.7	63.4±3.0	84.9±0.8	82.3±1.8	90.8±0.6	45.7±0.5	O.O.M.	30.5±2.5	21.9±3.0	29.0±1.3	64.5±0.5	56.9±0.6	62.1±0.3	8.4 (4.6)
H ² GCN	77.6±0.9	64.7±3.8	85.4±0.4	49.5±16.	75.8±11.	O.O.M.	O.O.M.	31.9±2.6	25.0±0.5	28.9±0.6	63.9±0.4	58.7±0.0	O.O.M.	8.9 (4.9)
APPNP	80.0±0.6	67.1±2.8	84.6±0.5	84.2±1.7	92.5±0.3	53.4±1.3	O.O.M.	30.9±4.7	23.9±3.2	26.1±1.0	63.7±0.9	47.3±0.3	57.4±0.4	7.6 (4.8)
GPR-GNN	78.8±1.3	64.2±4.0	85.1±0.7	85.0±1.0	92.6±0.3	58.5±0.8	O.O.M.	31.7±4.7	26.2±1.6	29.5±1.1	64.5±0.4	57.6±0.2	67.6±0.1	5.4 (3.7)
GAT	78.2±1.2	65.8±4.0	83.6±0.2	85.4±1.4	91.7±0.5	58.2±1.0	O.O.M.	39.1±4.1	28.6±0.6	26.4±0.4	60.5±0.8	O.O.M.	O.O.M.	7.5 (3.7)
SLIMG	77.8±1.1	67.1±2.3	84.6±0.5	86.3±0.7	91.8±0.5	66.3±0.3	84.9±0.0	40.8±3.2	31.1±0.7	30.9±0.6	68.2±0.6	59.7±0.1	73.9±0.1	1.9 (1.5)





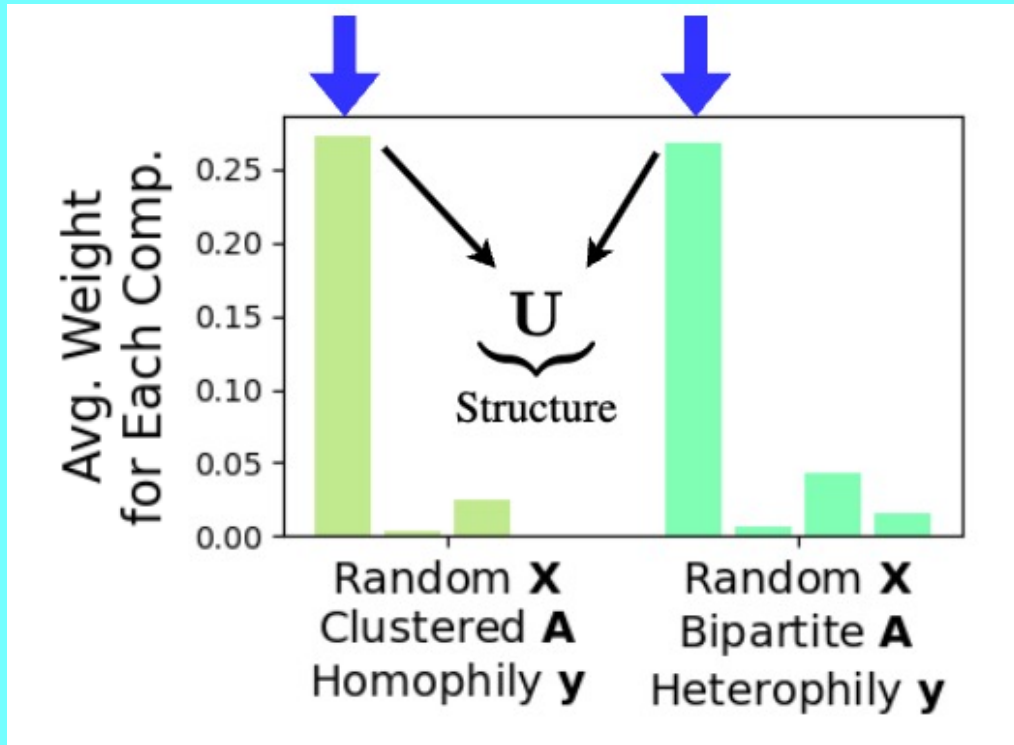
E2: Interpretable



structure
features
neighbors
grand-neighb.

$$\underbrace{\mathbf{U}}_{\text{Structure}} \parallel \underbrace{g(\mathbf{X})}_{\text{Node features}} \parallel \underbrace{g(\mathbf{A}_{\text{row}}^2 \mathbf{X})}_{\text{2-step neighbors}} \parallel \underbrace{g(\tilde{\mathbf{A}}_{\text{sym}}^2 \mathbf{X})}_{\text{Neighbors}}$$

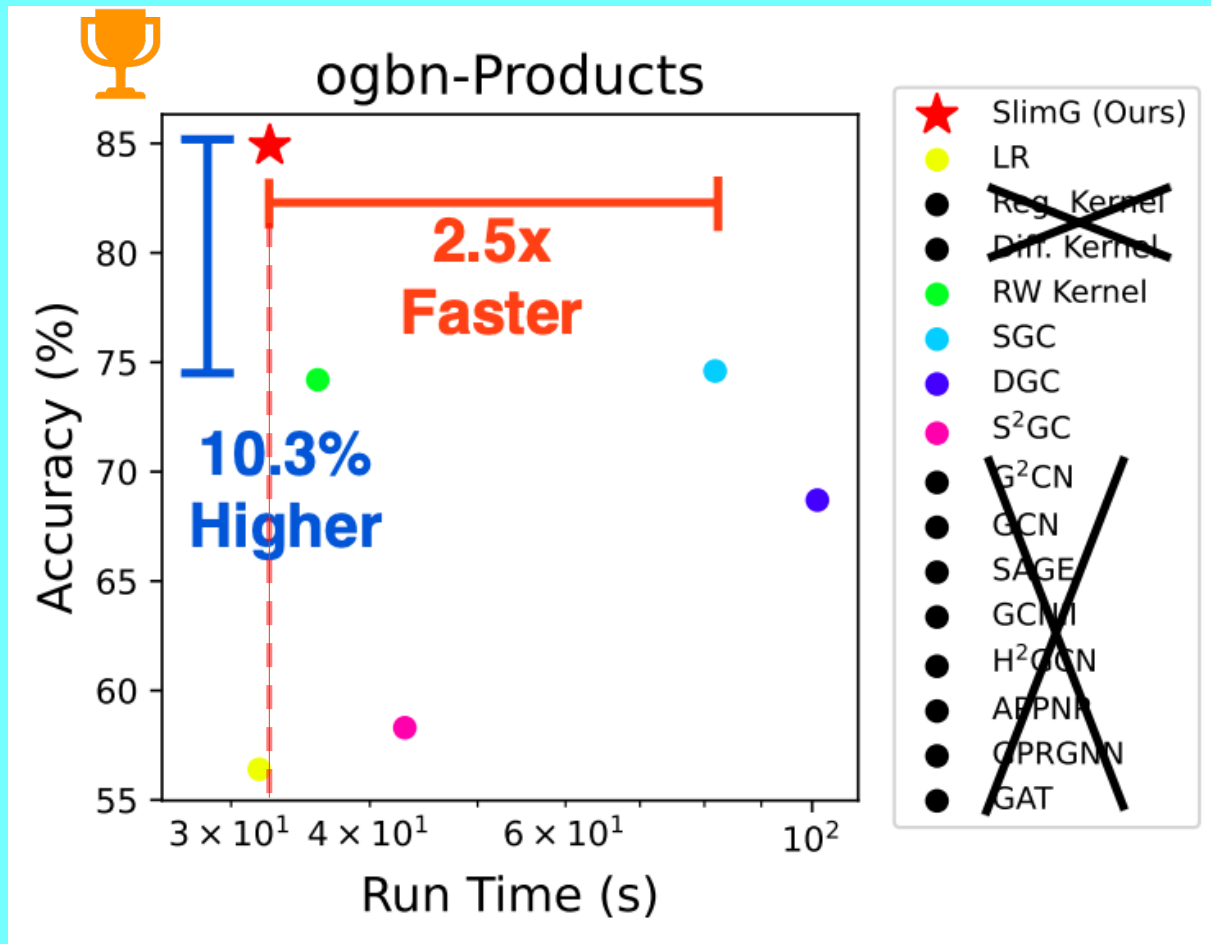
E2: Interpretable



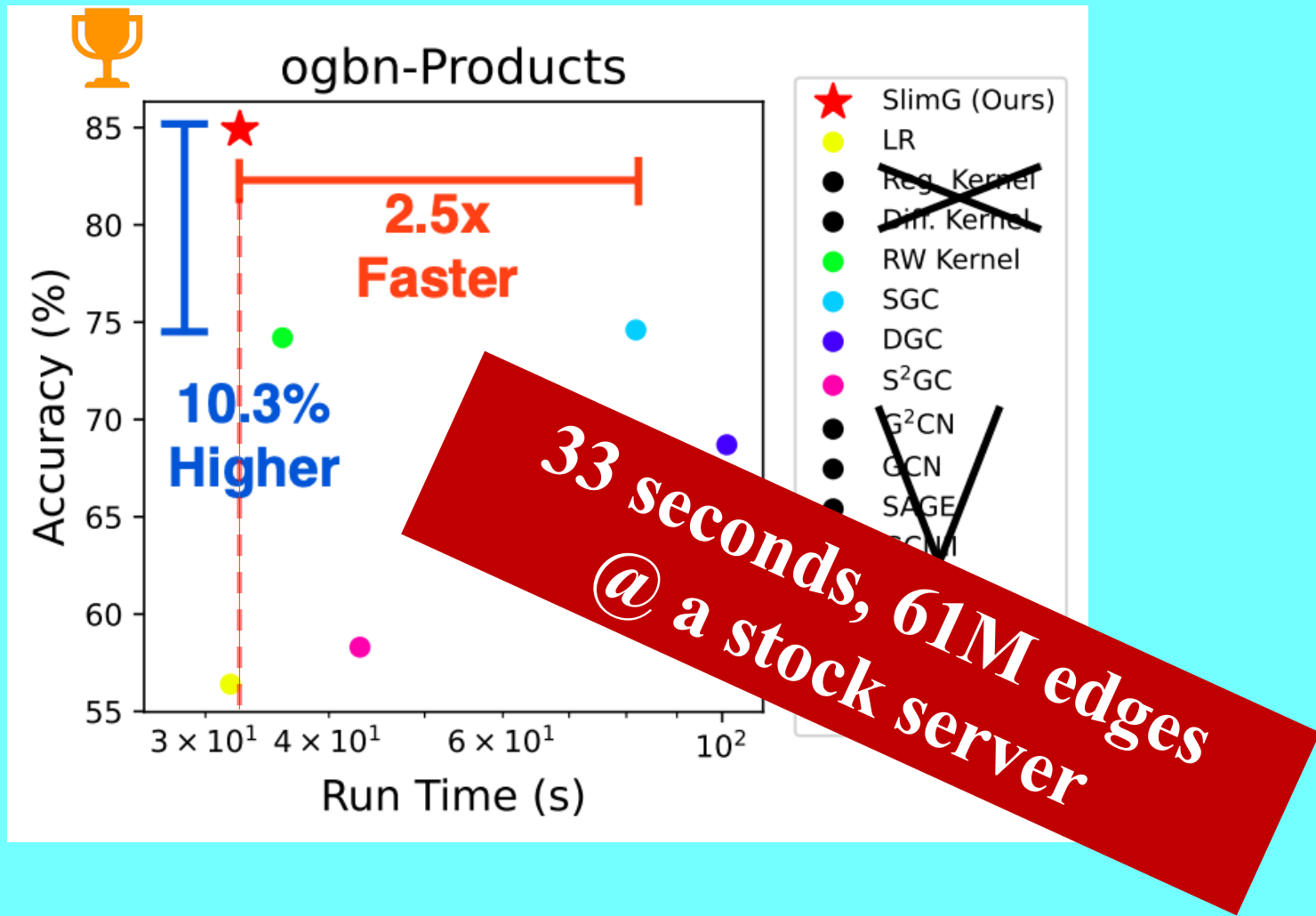
*structure
features
neighbors
grand-neighb.*

$$\underbrace{U}_{\text{Structure}} \parallel \underbrace{g(X)}_{\text{Node features}} \parallel \underbrace{g(A_{\text{row}}^2 X)}_{\text{2-step neighbors}} \parallel \underbrace{g(\tilde{A}_{\text{sym}}^2 X)}_{\text{Neighbors}}$$

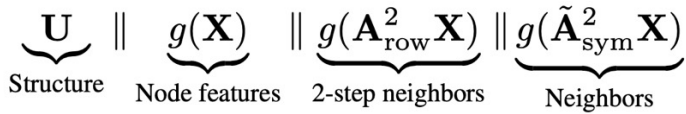
E3: Fast & Scalable



E3: Fast & Scalable



Conclusions



✓ Q1: Reasons

Model	Cora	CiteSeer	PubMed	Comp.	Photo	ArXiv	Products	Cham.	Squirrel	Actor	Penn94	Twitch	Pocec	Avg. Rank
LR	51.5±1.2	52.9±4.5	79.9±0.5	73.9±1.2	79.3±1.5	48.3±1.9	56.4±0.5	24.9±1.7	26.7±1.9	27.8±0.8	63.5±0.5	53.0±0.1	61.3±0.0	11.7 (4.2)
Reg. Kernel	67.8±2.5	62.1±4.4	83.4±1.4	80.3±1.4	87.1±1.2	O.O.M.	O.O.M.	29.4±2.6	24.3±2.3	29.6±1.4	O.O.M.	O.O.M.	O.O.M.	12.2 (3.8)
Diff. Kernel	70.6±1.5	62.7±3.8	82.1±0.4	83.1±1.0	89.8±0.6	O.O.M.	O.O.M.	34.5±7.9	28.3±1.5	24.7±0.9	53.5±0.8	O.O.M.	O.O.M.	11.8 (2.5)
RW Kernel	72.7±1.7	64.1±3.9	83.1±0.7	84.2±0.7	90.6±0.7	63.2±0.2	74.2±0.0	34.9±3.5	25.0±1.6	26.4±1.1	63.1±0.7	57.6±0.1	59.5±0.0	8.3 (3.3)
SGC	76.2±1.1	65.8±3.9	84.1±0.8	83.7±1.6	90.1±0.9	65.0±3.4	74.6±5.1	38.1±4.5	33.1±1.0	24.6±0.8	64.0±1.1	56.5±0.1	69.8±0.0	6.6 (4.2)
DGC	77.8±1.4	66.1±4.2	84.3±0.6	83.9±0.7	90.4±0.2	65.2±4.0	68.7±13.	37.2±3.7	29.2±1.2	25.2±2.1	62.5±0.4	58.2±0.2	60.7±0.1	6.6 (3.2)
S ² GC	78.3±1.5	66.9±4.4	84.3±0.3	83.1±0.8	90.1±0.8	62.0±7.4	58.3±18.	34.9±4.9	27.6±1.8	26.7±1.8	63.1±0.5	58.7±0.1	61.2±0.0	6.6 (2.7)
G ² CN	76.6±1.5	64.2±3.3	81.4±0.6	82.8±1.6	88.8±0.5	O.O.M.	O.O.M.	40.7±2.9	32.1±1.5	24.3±0.5	O.O.M.	O.O.M.	O.O.M.	10.5 (4.5)
GCN	76.0±1.2	65.0±2.9	84.3±0.5	85.1±0.9	91.6±0.5	62.8±0.6	O.O.M.	38.5±3.0	31.4±1.8	26.8±0.4	62.9±0.7	57.0±0.1	63.9±0.4	6.3 (2.4)
SAGE	74.6±1.3	63.7±3.6	82.9±0.4	83.8±0.5	90.6±0.5	61.5±0.6	O.O.M.	39.8±4.3	27.0±1.3	27.8±0.9	O.O.M.	56.6±0.4	68.9±0.1	8.5 (3.5)
GCNII	77.8±1.7	63.4±3.0	84.9±0.8	82.3±1.8	90.8±0.6	45.7±0.5	O.O.M.	30.5±2.5	21.9±3.0	29.0±1.3	64.5±0.5	56.9±0.6	62.1±0.3	8.4 (4.6)
H ² GCN	77.6±0.9	64.7±3.8	85.4±0.4	49.5±1.6	75.8±11.	O.O.M.	O.O.M.	31.9±2.6	25.0±0.5	28.9±0.6	63.9±0.4	58.7±0.0	O.O.M.	8.9 (4.9)
APNP	80.0±0.6	67.1±2.3	84.6±0.5	84.2±1.7	92.5±0.3	53.4±1.3	O.O.M.	30.9±4.7	23.9±3.2	26.1±1.0	63.7±0.9	47.3±0.3	57.4±0.4	7.6 (4.8)
GPR-GNN	78.8±1.3	64.2±4.0	85.1±0.7	85.0±1.0	92.6±0.3	58.5±0.8	O.O.M.	31.7±4.7	26.2±1.6	29.5±1.1	64.5±0.4	57.6±0.2	67.6±0.1	5.4 (3.7)
GAT	78.2±1.2	65.8±4.0	83.6±0.2	85.4±1.4	91.7±0.5	58.2±1.0	O.O.M.	39.1±4.1	28.6±0.6	26.4±0.4	60.5±0.8	O.O.M.	O.O.M.	7.5 (3.7)
SLIMG	77.8±1.1	67.1±2.3	84.6±0.5	86.3±0.7	91.8±0.5	66.3±0.3	84.9±0.0	40.8±3.2	31.1±0.7	30.9±0.6	68.2±0.6	59.7±0.1	73.9±0.1	1.9 (1.5)

✓ Q2: Method

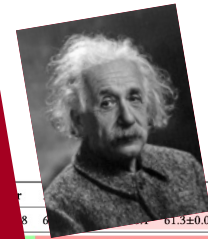


– Accurate, Robust, Interpretable, Fast, Scalable

✓ Q3: Sanity Checks



Conclusions



$$\underbrace{U}_{\text{Structure}} \parallel \underbrace{g(\mathbf{X})}_{\text{Node}} \parallel \underbrace{g(\mathbf{A}_{\text{row}}^2 \mathbf{X})}_{\text{Edge}} \parallel \underbrace{g(\tilde{\mathbf{A}})}_{\text{Graph}}$$

“Careful Simplicity”

Method	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	Avg. Rank
GCN	76.6±1.5	64.2±3.3	81.4±0.6	82.8±1.6	88.8±0.5	O.O.M.	O.O.M.	40.7±2.9	32.1±1.5	24.3±0.5	O.O.M.	O.O.M.	O.O.M.	O.O.M.	O.O.M.	O.O.M.	O.O.M.	O.O.M.	O.O.M.	O.O.M.	11.7 (4.2)
SAGE	74.6±1.3	63.7±3.6	82.9±0.4	83.8±0.5	90.6±0.5	61.5±0.6	O.O.M.	39.8±4.3	27.0±1.3	27.8±0.9	O.O.M.	O.O.M.	56.6±0.4	68.9±0.1	8.5 (3.5)						
GCNII	77.8±1.7	63.4±3.0	84.9±0.8	82.3±1.8	90.8±0.6	45.7±0.5	O.O.M.	30.5±2.5	21.9±3.0	29.0±1.3	64.5±0.5	56.9±0.6	62.1±0.3	8.4 (4.6)							
H ² GCN	77.6±0.9	64.7±3.8	85.4±0.4	49.5±1.6	75.8±1.1	O.O.M.	O.O.M.	31.9±2.6	25.0±0.5	28.9±0.6	63.9±0.4	58.7±0.0	O.O.M.	8.9 (4.9)							
APPNP	80.0±0.6	67.1±2.3	84.6±0.5	84.2±1.7	92.5±0.3	53.4±1.3	O.O.M.	30.9±4.7	23.9±3.2	26.1±1.0	63.7±0.9	47.3±0.3	57.4±0.4	7.6 (4.8)							
GPR-GNN	78.8±1.3	64.2±4.0	85.1±0.7	85.0±1.0	92.6±0.3	58.5±0.8	O.O.M.	31.7±4.7	26.2±1.6	29.5±1.1	64.5±0.4	57.6±0.2	67.6±0.1	5.4 (3.7)							
GAT	78.2±1.2	65.8±4.0	83.6±0.2	85.4±1.4	91.7±0.5	58.2±1.0	O.O.M.	39.1±4.1	28.6±0.6	26.4±0.4	60.5±0.8	O.O.M.	O.O.M.	7.5 (3.7)							
SLIMG	77.8±1.1	67.1±2.3	84.6±0.5	86.3±0.7	91.8±0.5	66.3±0.3	84.9±0.0	40.8±3.2	31.1±0.7	30.9±0.6	68.2±0.6	59.7±0.1	73.9±0.1	1.9 (1.5)							



Q2: Method



– Accurate, Robust, Interpretable, Fast, Scalable



Q3: Sanity Checks



Thank You!



Jaemin
 YOO*



Meng-Chieh
 (Jeremy) LEE*



Shubhranshu
 SHEKHAR

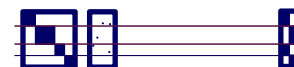


Christos
 FALOUTSOS

✓Q1: Reasons

✓Q2: Method

✓Q3: Sanity Checks



Model	Cora	CiteSeer	PubMed	Comp.	Photo	ArXiv	Products	Cham.	Squirrel	Actor	Penn94	Twitch	Pokec	Avg. Rank
LR	51.5±1.2	52.9±4.5	79.9±0.5	73.9±1.2	79.3±1.5	48.3±1.9	56.4±0.5	24.9±1.7	26.7±1.9	27.8±0.8	63.5±0.5	53.0±0.1	61.3±0.0	11.7 (4.2)
Reg. Kernel	67.8±2.5	62.1±4.4	83.4±1.4	80.3±1.4	87.1±1.2	O.O.M.	O.O.M.	29.4±2.6	24.3±2.3	29.6±1.4	O.O.M.	O.O.M.	O.O.M.	12.2 (3.8)
Diff. Kernel	70.6±1.5	62.7±3.8	82.1±0.4	83.1±1.0	89.8±0.6	O.O.M.	O.O.M.	34.5±7.9	28.3±1.5	24.7±0.9	33.5±0.9	O.O.M.	O.O.M.	11.8 (2.5)
RW Kernel	72.7±1.7	64.1±1.9	83.1±0.7	84.2±0.7	90.6±0.7	63.2±0.2	74.2±0.0	34.9±3.5	25.0±1.6	26.4±1.1	53.1±0.7	57.6±0.1	59.3±0.0	8.3 (3.3)
SGC	76.2±1.1	65.8±3.9	84.1±0.5	83.7±1.6	90.3±0.9	65.0±3.4	74.6±5.1	38.1±4.5	27.7±1.0	24.6±0.8	44.0±1.1	56.5±0.1	69.8±0.0	6.6 (4.2)
DOC	77.8±1.4	66.1±4.2	84.3±0.6	83.9±0.7	90.4±0.2	65.2±4.0	68.7±1.3	37.2±3.7	29.2±1.2	25.2±2.1	62.5±0.4	58.2±0.2	60.7±0.1	6.6 (2.2)
S ² GC	78.3±1.5	66.9±4.4	84.3±0.3	83.1±0.8	90.1±0.8	62.0±7.4	58.3±1.8	34.9±4.9	27.6±1.8	26.7±1.8	63.1±0.5	58.7±0.1	61.2±0.0	6.6 (2.7)
G ² CN	76.6±1.5	64.2±3.3	81.4±0.6	82.8±1.6	88.8±0.5	O.O.M.	O.O.M.	40.7±2.9	32.1±1.5	24.3±0.5	O.O.M.	O.O.M.	O.O.M.	10.5 (4.5)
GCN	76.0±1.2	65.0±2.9	84.3±0.5	85.1±0.9	91.6±0.5	62.8±0.6	O.O.M.	38.5±3.0	31.4±1.8	26.8±0.4	62.9±0.7	57.0±0.1	63.9±0.4	6.3 (2.4)
SAGE	77.6±1.3	63.7±3.6	82.9±0.4	83.8±0.5	90.6±0.5	61.2±0.6	O.O.M.	39.8±4.3	27.0±1.3	27.8±0.9	O.O.M.	56.6±0.4	68.9±0.1	8.3 (3.5)
GCNII	77.8±1.7	63.4±3.0	84.9±0.8	82.9±1.8	90.8±0.6	65.7±0.5	O.O.M.	39.5±2.5	21.9±1.0	29.0±1.3	64.5±0.5	56.9±0.6	62.1±0.3	8.4 (4.6)
HFGCN	77.6±0.9	64.7±3.8	85.4±0.5	85.4±0.5	89.5±1.6	75.8±1.1	O.O.M.	31.9±2.6	25.0±0.5	28.9±0.6	63.9±0.4	58.7±0.0	O.O.M.	8.9 (4.8)
APFNP	80.0±0.6	70.1±2.8	84.6±0.5	84.2±1.7	92.5±0.3	53.4±1.3	O.O.M.	30.9±8.7	23.9±3.2	26.1±1.0	63.7±0.9	47.3±0.3	57.4±0.4	7.6 (4.8)
GPR-GNN	78.8±1.3	64.2±4.0	85.1±0.7	85.0±1.0	90.9±0.9	58.5±0.8	O.O.M.	31.7±4.7	26.2±1.6	29.5±1.1	64.5±0.4	57.6±0.2	67.6±0.1	5.4 (3.7)
GAT	79.2±1.2	63.8±4.0	83.6±0.2	86.6±0.6	91.7±0.5	58.2±1.0	O.O.M.	39.1±4.1	28.6±0.6	26.4±0.4	69.5±0.4	O.O.M.	O.O.M.	7.5 (3.7)
SLiNG	77.8±1.1	67.1±2.3	84.6±0.5	86.3±0.7	91.8±0.5	66.3±0.3	64.9±0.0	41.3±2.1	31.1±0.7	30.2±0.2	63.2±0.6	59.2±0.1	73.9±0.1	7.9 (4.1)

