

Are Graph Convolutional Networks Fully Exploiting Graph Structure?

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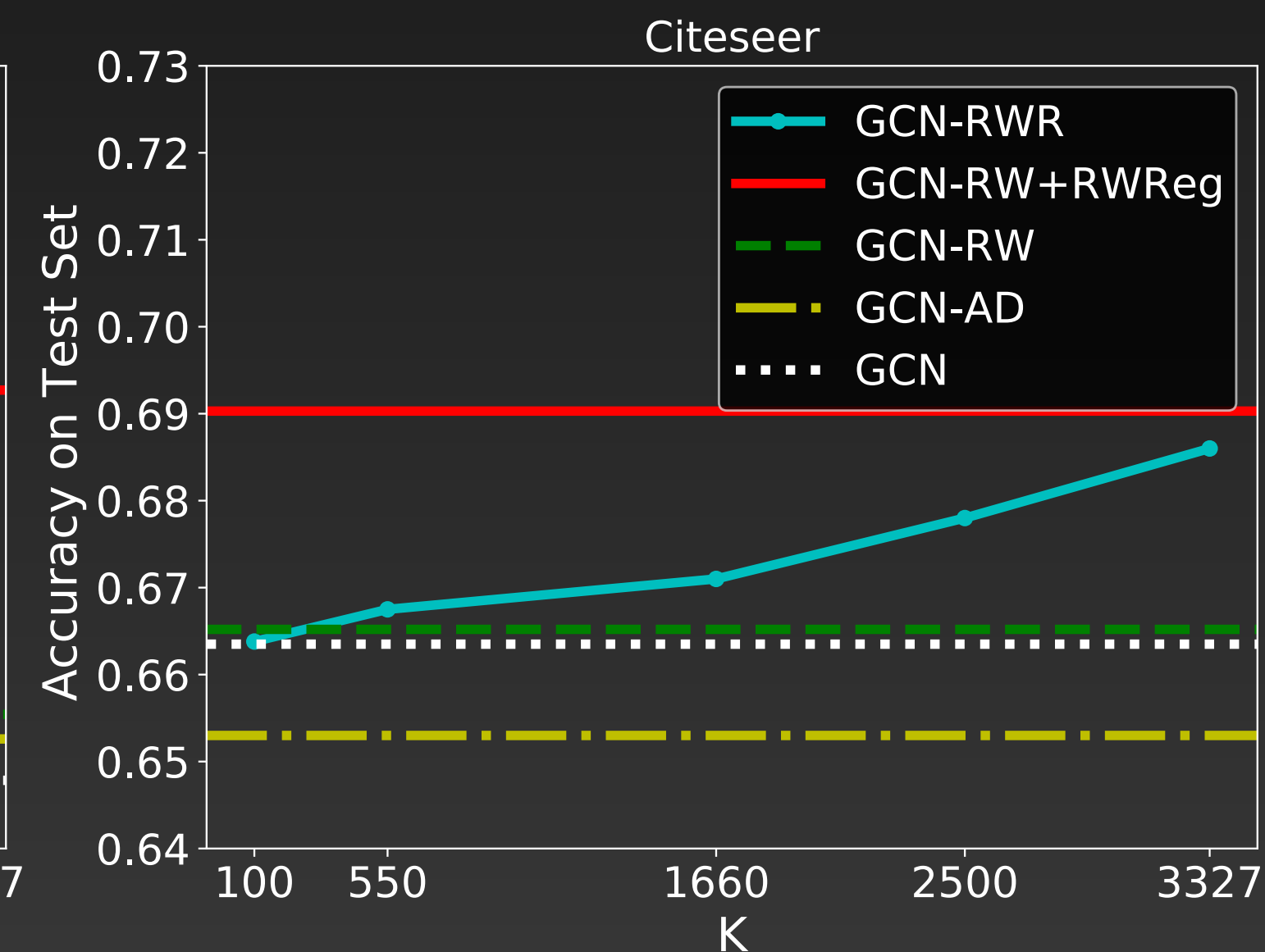
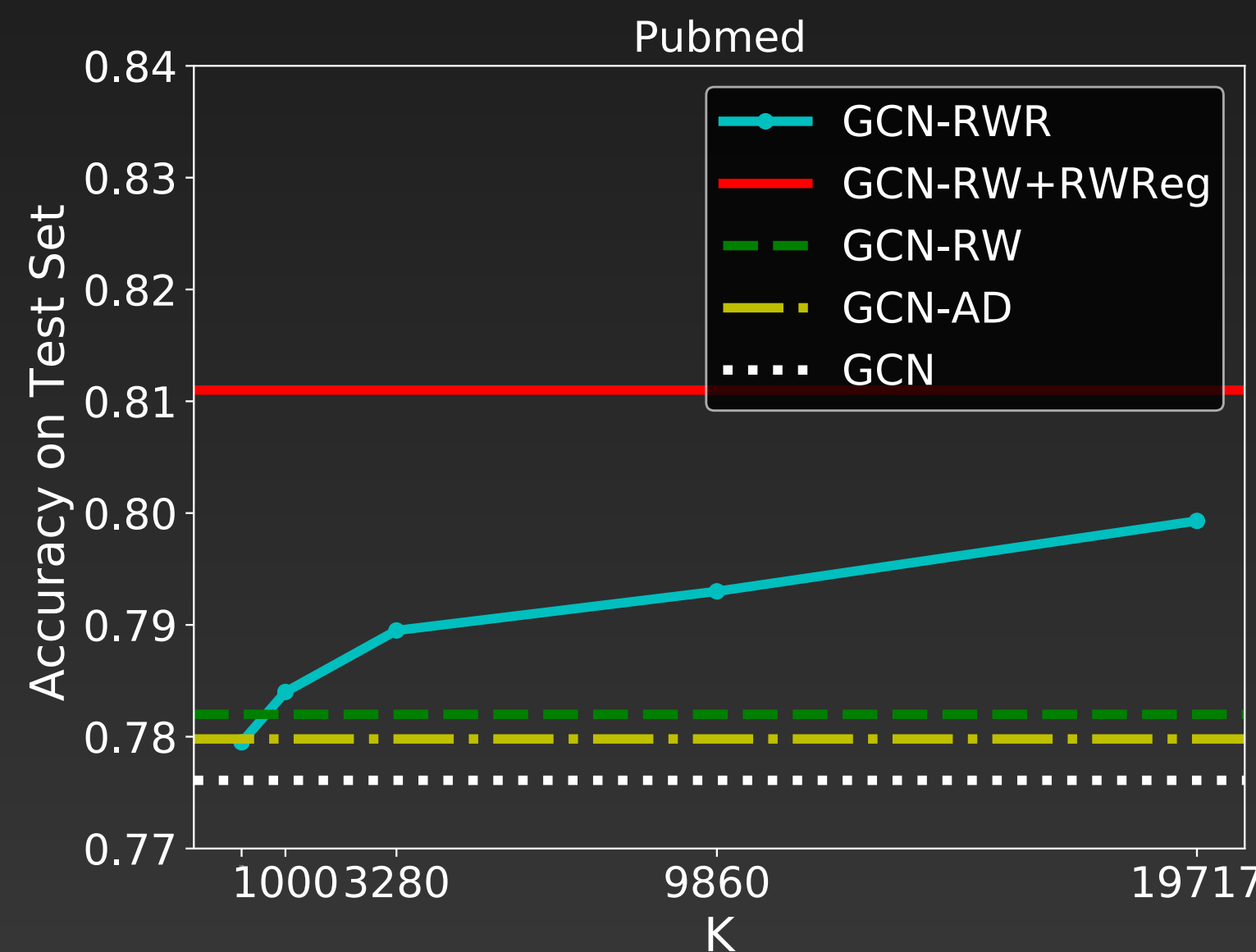
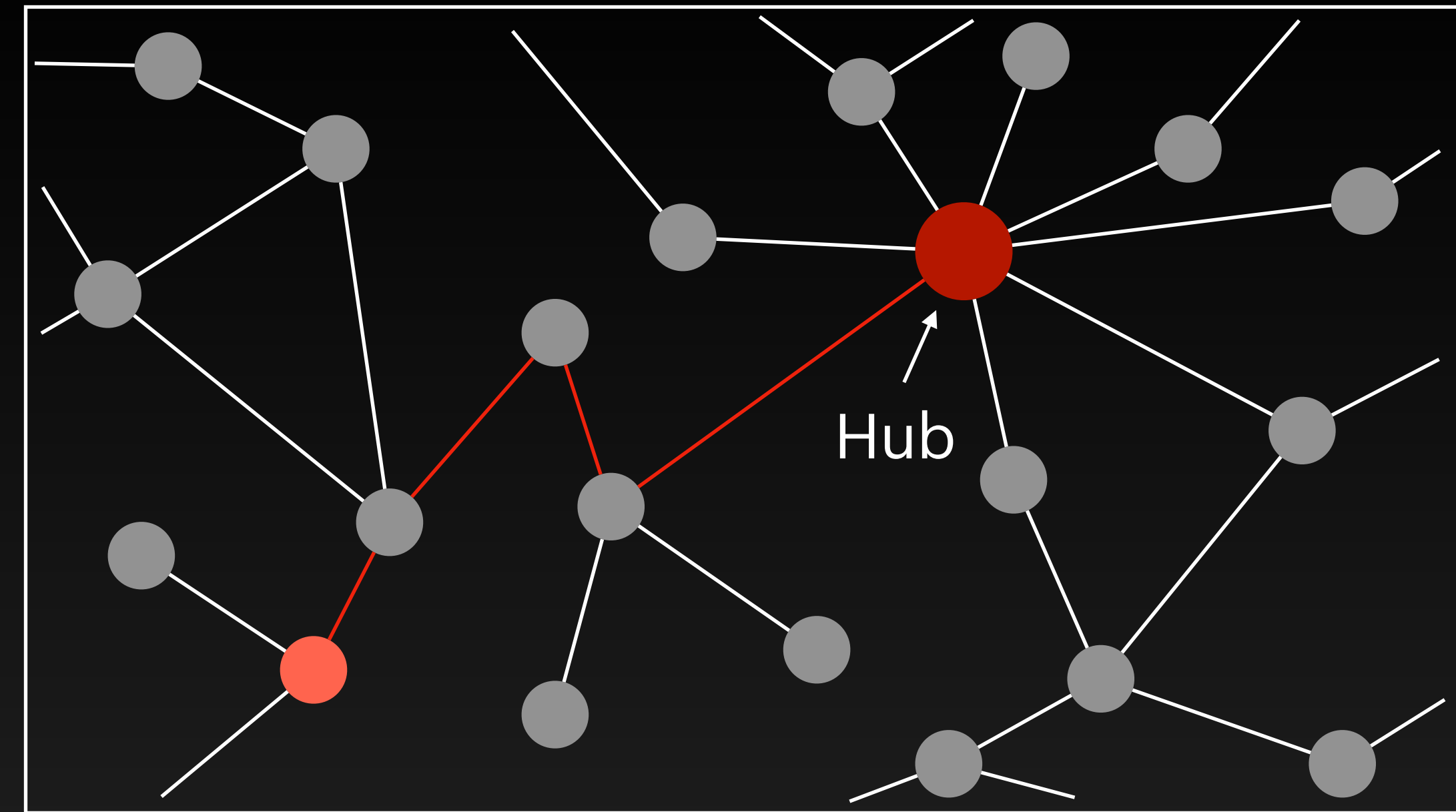
What is the impact of missing long range dependencies?

Three levels of structural knowledge injection:

- Adjacency Matrix **[AD]** $\hat{X} = [X, A]$
- RWR Statistics **[RW]** $\hat{X} = [X, S]$
- RWR Statistics + Regularization Term **[RW+RWR]**
 $\hat{X} = [X, S] + \mathcal{L}_{\text{RWReg}} = \sum_{i,j} s_{i,j} ||H_{i,:} - H_{j,:}||^2$

Why Random Walks with Restart?

- Empirical Results
RWR capture non-trivial dependencies between nodes
- Theoretical Results
RWR-based colourings speed up the WL algorithm



RWR Regularization **[RWR]** 📈 >5% accuracy

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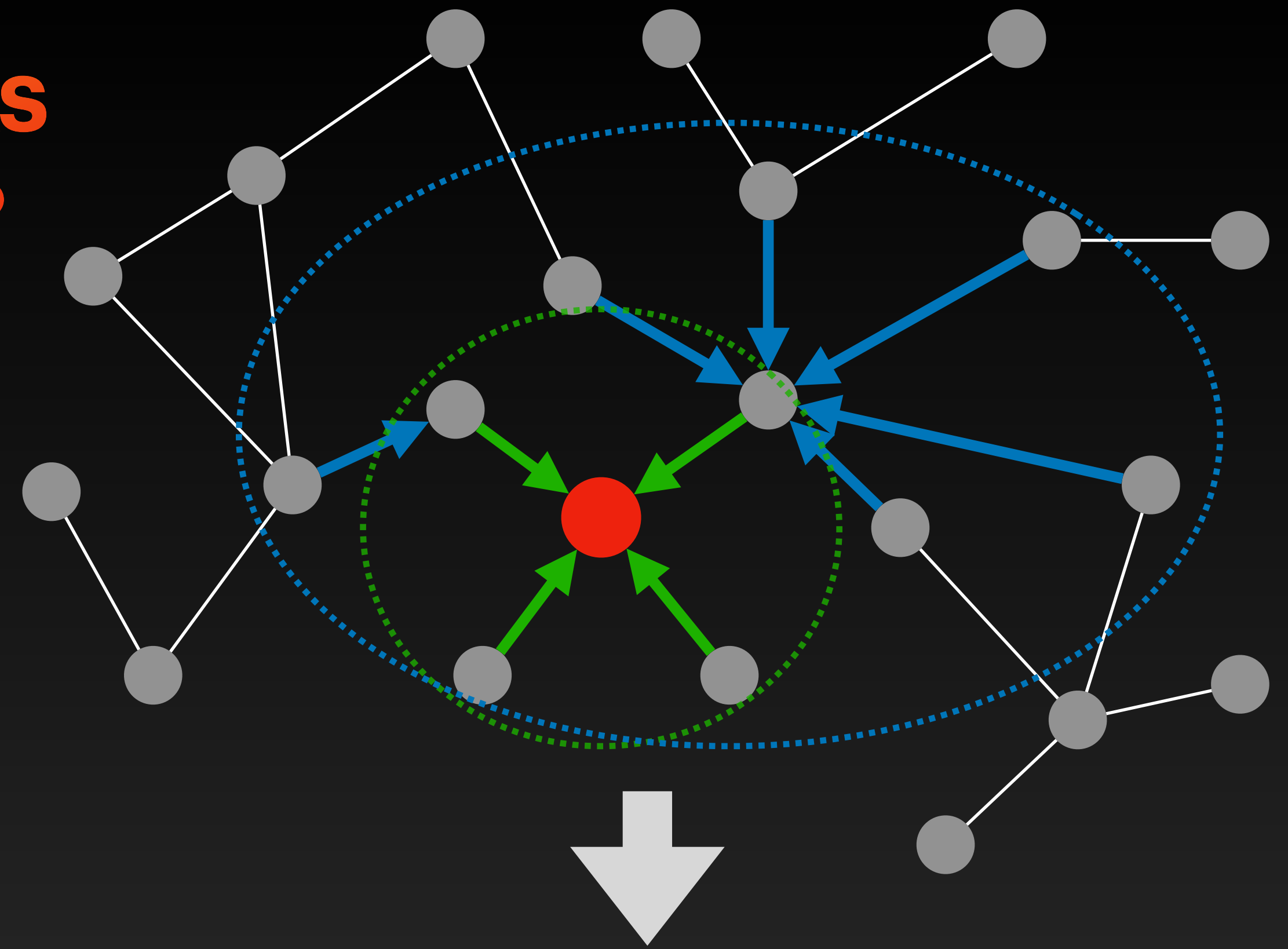
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Graph Convolutional Networks

Message Passing Iteration l :

- Create Messages $m_{ij}^l = MSG(h_i^{l-1}, h_j^{l-1}, e_{ij})$
- Aggregate messages from neighbours $M_i^l = AGG(\{m_{ij}^l | v_j \in N(v_i)\})$
- Update node representation $h_i^l = UPDATE(\{M_i^l, h_i^{l-1}\})$

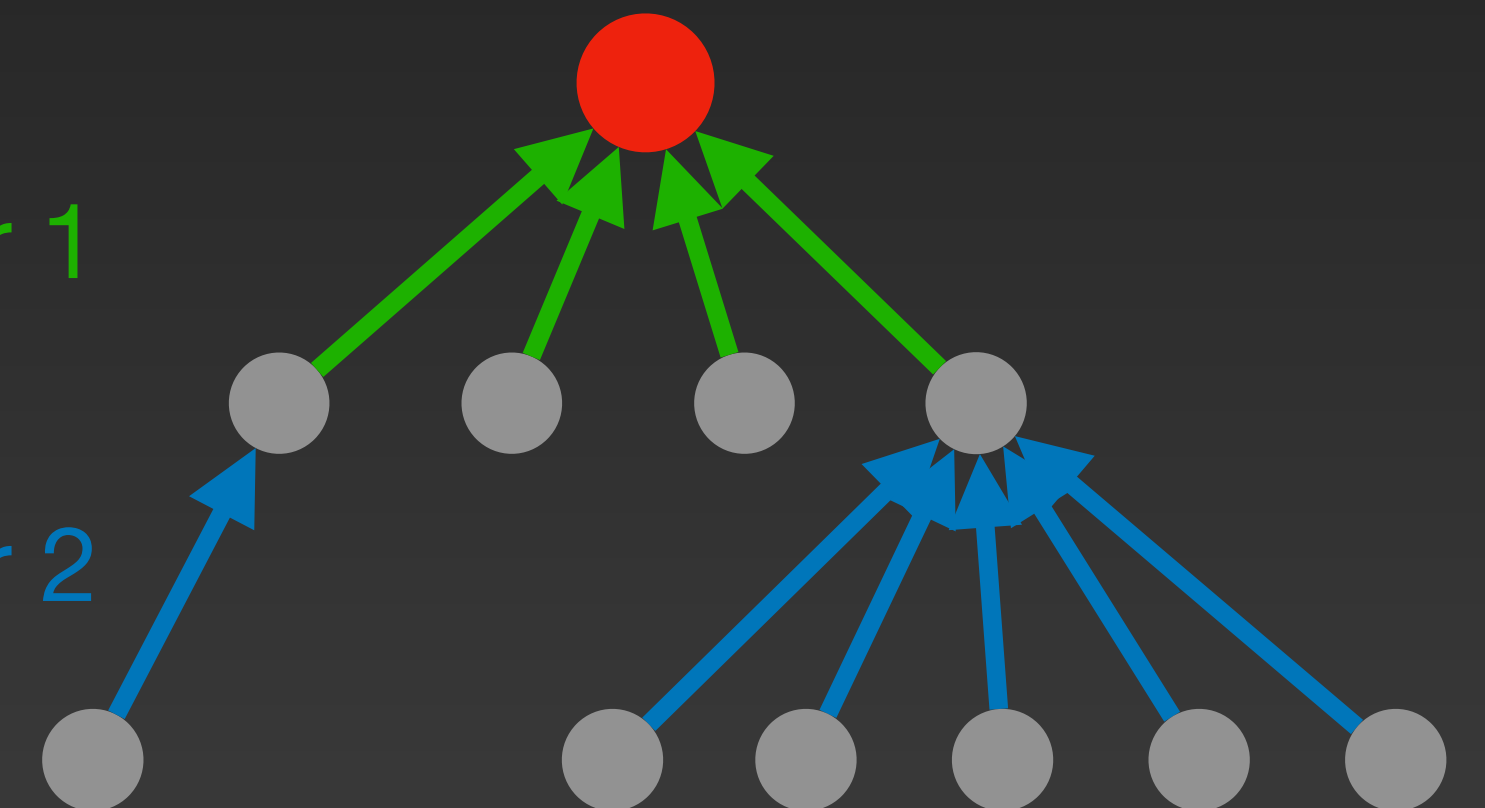
After k message passing iterations a node will access information from its k -hop neighbourhood



One computation graph per node

Message Passing Layer 1

Message Passing Layer 2



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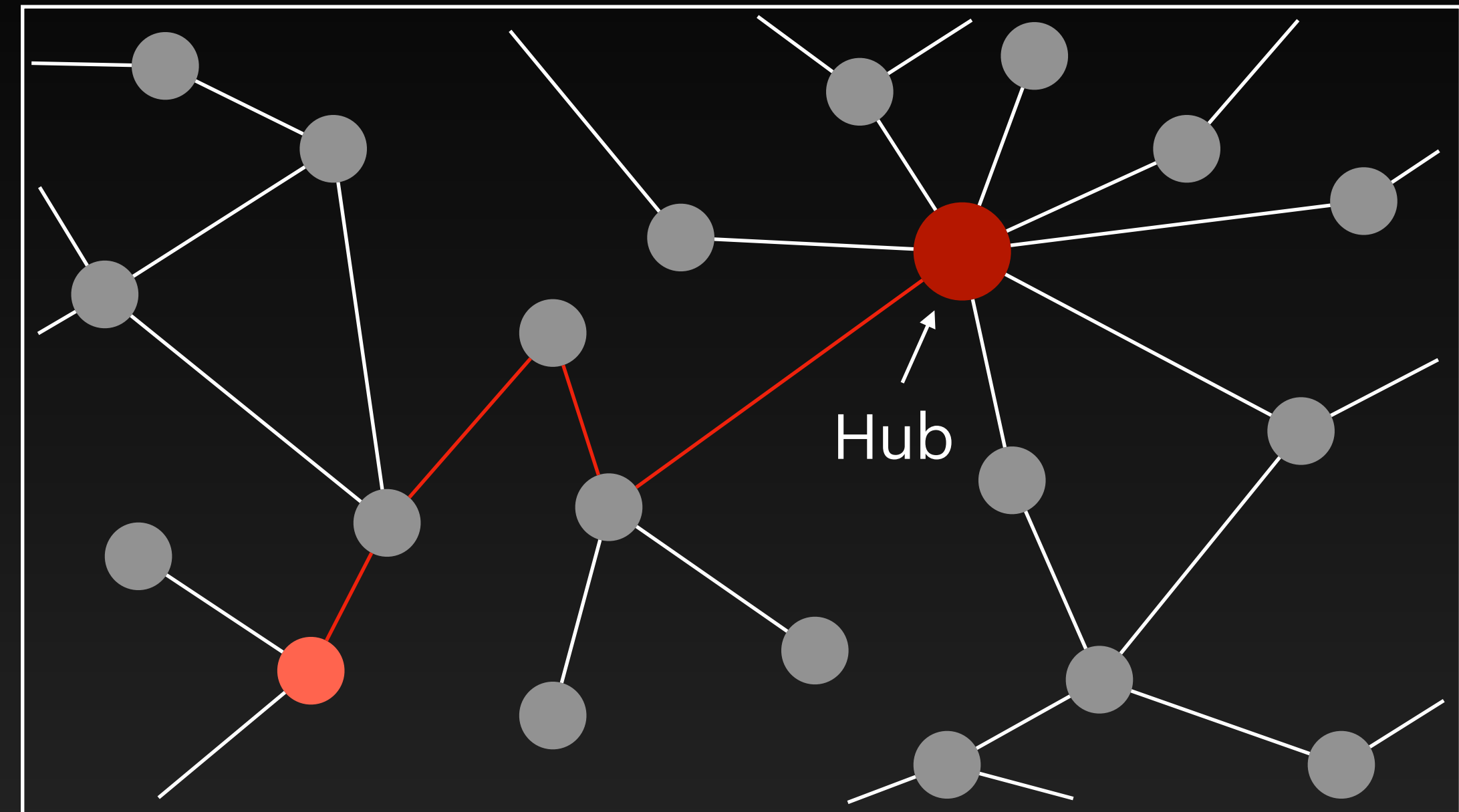
Motivation:

Information in a graph is not limited to local neighbourhoods, however oversmoothing is preventing GCNs from long-range communications.

How much information is lost?

Can we try to quantify it?

How can we introduce long-range dependencies?



Quantifying the impact of missing long range dependencies

Three levels of structural knowledge injection:

- Adjacency Matrix $\hat{X} = [X, A]$
- RWR Statistics $\hat{X} = [X, S]$
- RWR Statistics + Regularization Term

$$\hat{X} = [X, S] + \mathcal{L}_{\text{RWReg}} = \sum_{i,j} s_{i,j} ||H_{i,:} - H_{j,:}||^2$$

Empirical Evaluation



5 architectures:

- GCN
- GAT
- GraphSage
- DiffPool
- k-GNN

3 tasks:

- Node Classification
- Graph Classification
- Triangle Count

Results:

-  Accuracy
-  Generalization on Inductive Tasks
- All Architectures are affected

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Why Random Walks with Restart?

Theoretical Results

RWR-based colourings speed up the WL algorithm

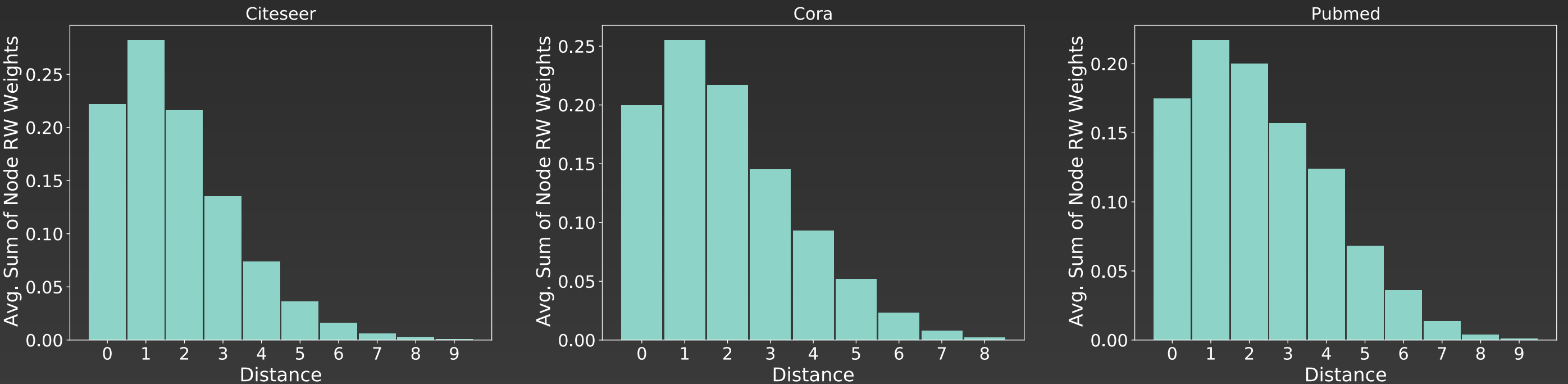
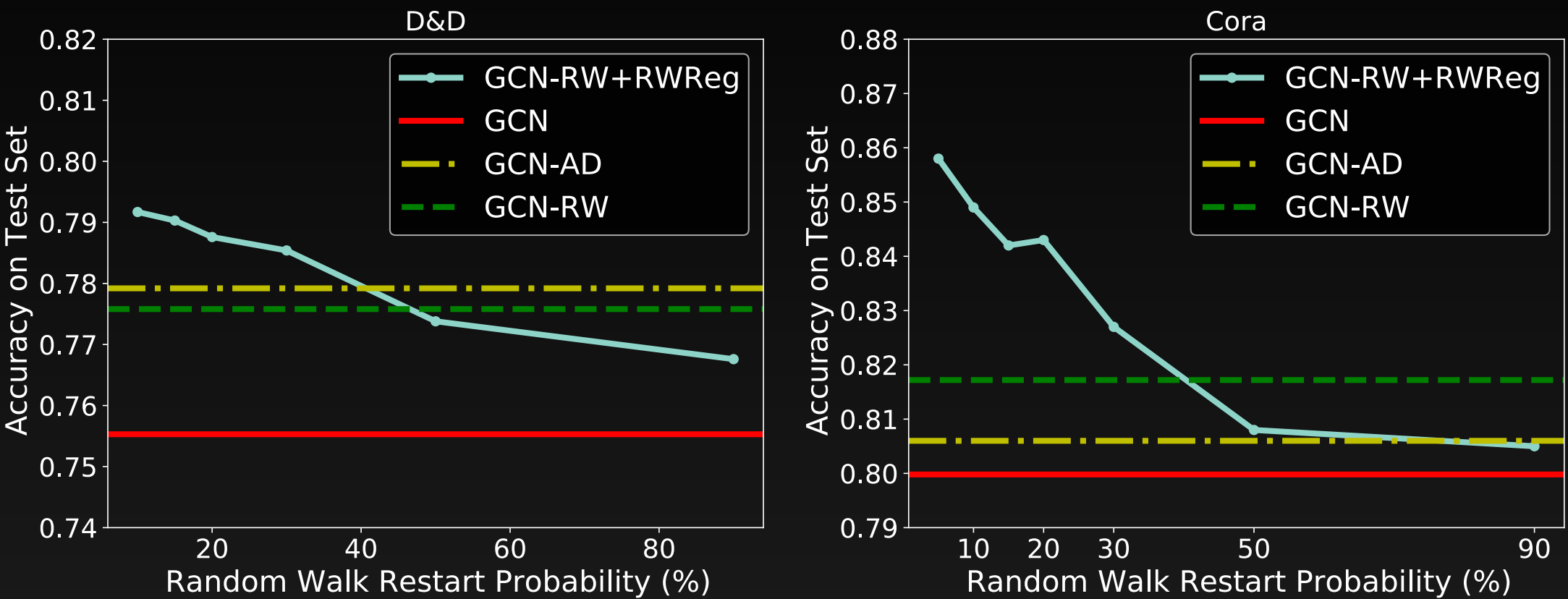
Proposition Let $G_1 = (V_1, E_1)$ and $G_2 = (V_2, E_2)$ be two non-isomorphic graphs for which the 1-WL algorithm terminates with the correct answer after k iterations and starting from the labelling of all 1's. Then the k -long random walk representations of G_1 and G_2 are different.

Empirical Results

RWR capture non-trivial dependencies between nodes

Dataset	Average Kendall Tau-b
Cora	0.729 ± 0.082
Pubmed	0.631 ± 0.057
Citeseer	0.722 ± 0.171

Changing Restart Probability




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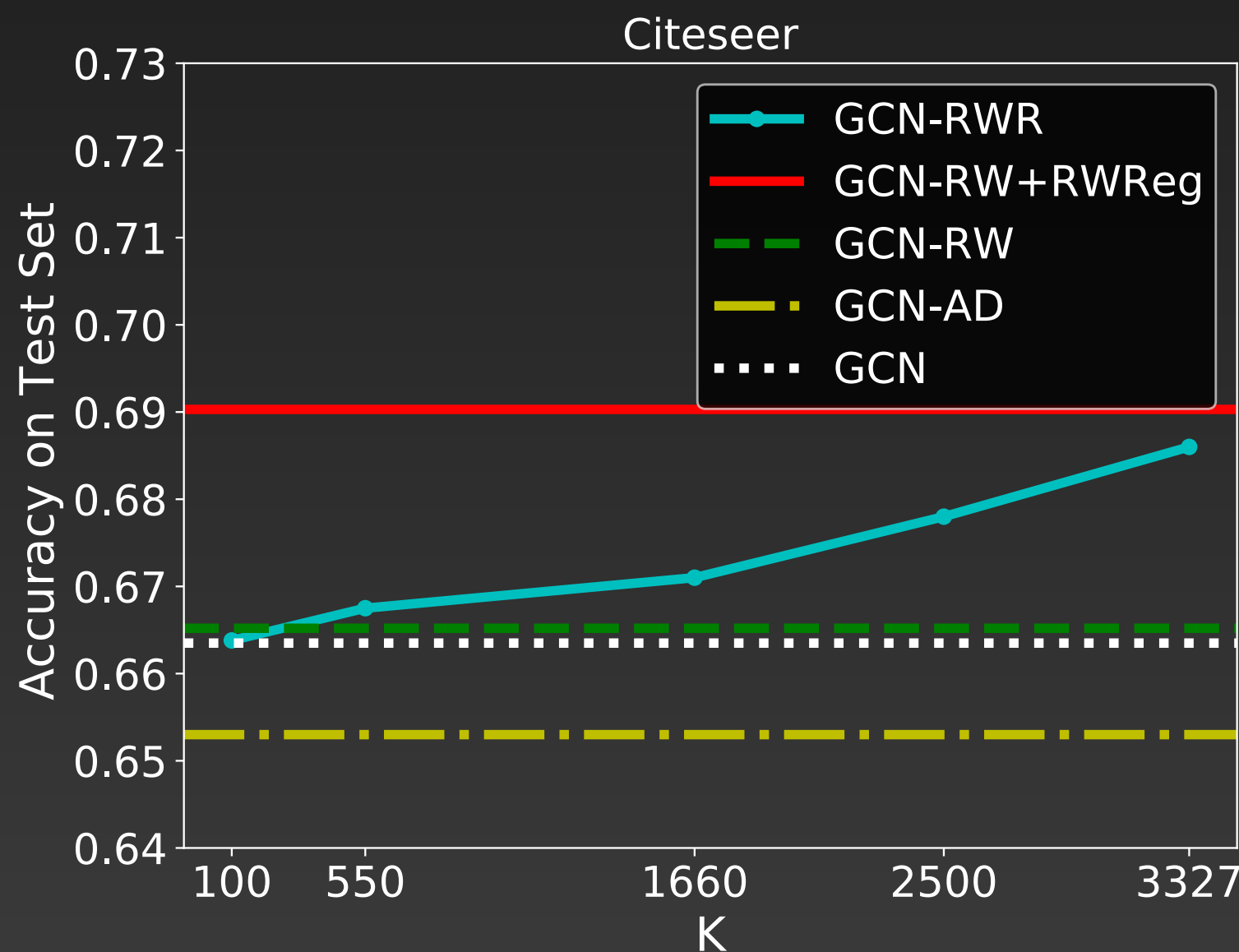
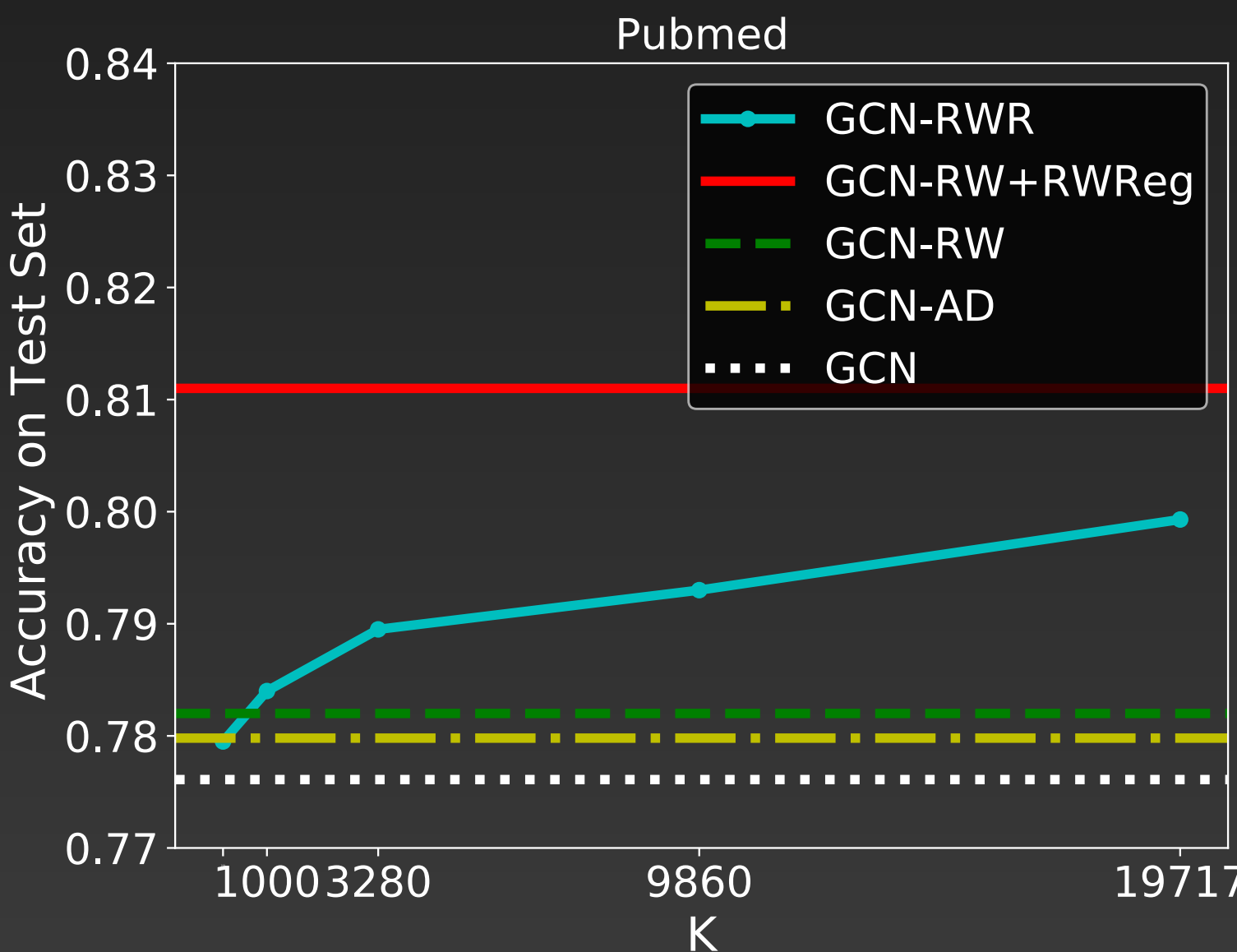
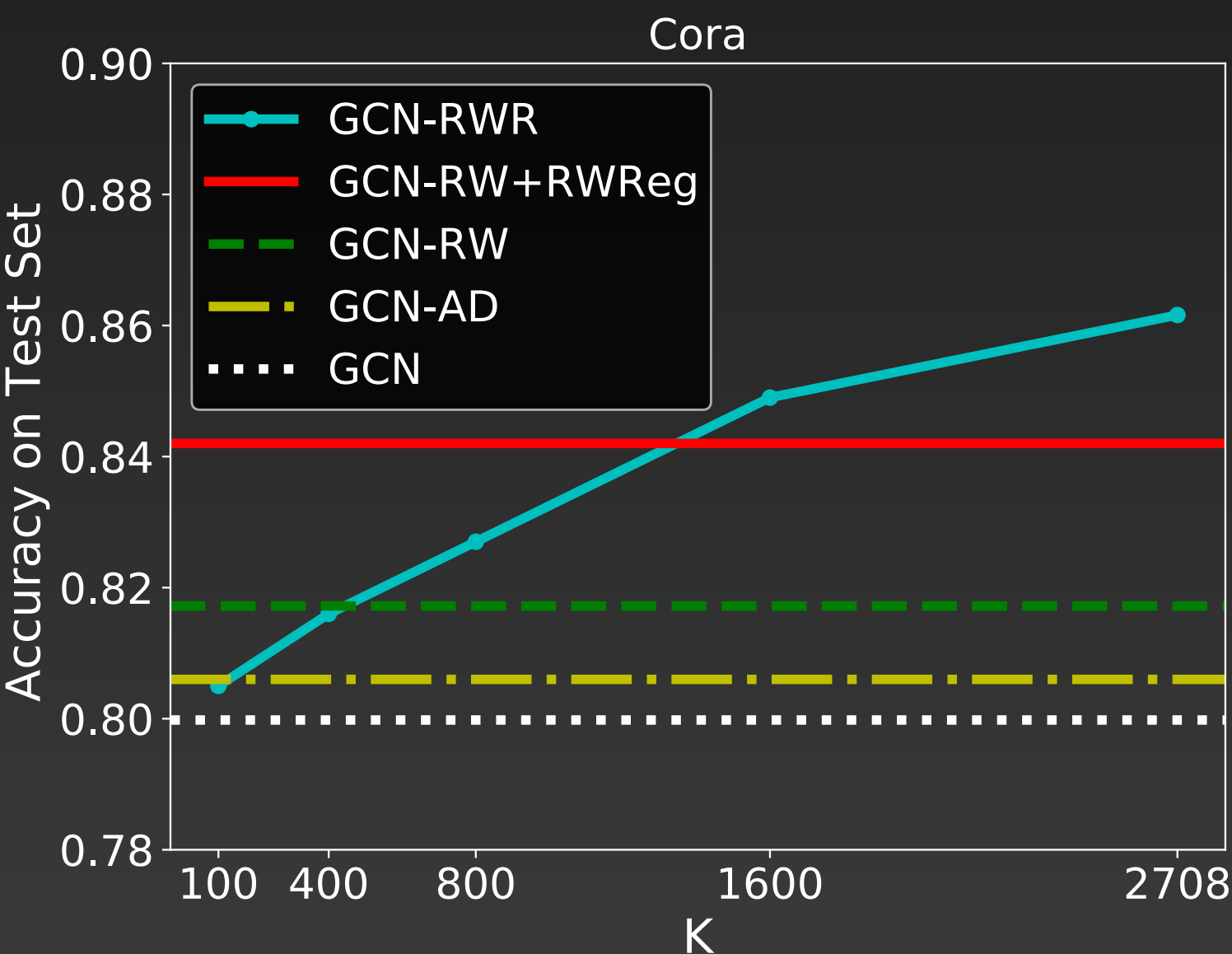
RWReg: a Tractable Method

[RWR] $\mathcal{L}_{\text{RWReg}} = \sum_{i,j} s_{i,j} ||H_{i,:} - H_{j,:}||^2$

- Permutation Invariant
- No additional operations at inference time
- No additional features
-  > 5% accuracy improvement
- No additional parameters
- Control scalability with K

Results on Node Classification (Accuracy)

	GCN	GCN RW+RWReg	GCN RWR
Cora	0.7998 ±0.029	0.8420 ±0.026	0.8616 ±0.025
Pubmed	0.7761 ±0.022	0.8110 ±0.037	0.7993 ±0.034
Citeseer	0.6635 ±0.095	0.6903 ±0.102	0.6860 ±0.096



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Empirical Results

The impact of structural knowledge:

- Improvements on all architectures
- Improvements on all tasks
- RW & RW+RWReg give important contributions
- Graph Classification is more affected
- Increase in generalization in Triangle Counting

Triangle Count (MSE)

Model	TRIANGLES Test Set		
	Global	Small	Large
GCN	2.2900	1.3111	3.6082
GCN-AD	4.7469	1.1628	5.9717
GCN-RW	2.0449	1.1008	2.9889
GCN-RW+RWReg	2.0298	1.1664	2.8932

Node Classification (Accuracy)

Model	Dataset	Structural Information			
		none	AD	RW	RW+RWReg
GCN	Cora	0.7998 ± 0.029	0.8060 ± 0.035	0.8172 ± 0.025	0.8420 ± 0.026
	Pubmed	0.7761 ± 0.022	0.7798 ± 0.070	0.7820 ± 0.042	0.8110 ± 0.037
	Citeseer	0.6635 ± 0.095	0.6530 ± 0.104	0.6652 ± 0.098	0.6903 ± 0.102
GraphSage	Cora	0.8061 ± 0.017	0.8031 ± 0.014	0.8067 ± 0.014	0.8270 ± 0.015
	Pubmed	0.8071 ± 0.016	0.8037 ± 0.013	0.8076 ± 0.015	0.8202 ± 0.010
	Citeseer	0.6811 ± 0.021	0.6889 ± 0.020	0.6930 ± 0.019	0.7280 ± 0.020
GAT	Cora	0.8150 ± 0.021	0.8236 ± 0.019	0.8332 ± 0.020	0.8480 ± 0.019
	Pubmed	0.8046 ± 0.011	0.7966 ± 0.014	0.8114 ± 0.009	0.8287 ± 0.010
	Citeseer	0.6646 ± 0.008	0.6720 ± 0.017	0.6866 ± 0.009	0.7011 ± 0.011

Graph Classification (Accuracy)

Model	Dataset	Structural Information			
		none	AD	RW	RW+RWReg
GCN	ENZYMES	0.5702 ± 0.052	0.5916 ± 0.076	0.5845 ± 0.055	0.6166 ± 0.065
	D&D	0.7553 ± 0.028	0.7792 ± 0.022	0.7758 ± 0.023	0.7903 ± 0.023
	PROTEINS	0.7400 ± 0.035	0.7758 ± 0.042	0.7848 ± 0.034	0.7957 ± 0.032
DiffPool	ENZYMES	0.6610 ± 0.031	0.7113 ± 0.027	0.6876 ± 0.025	0.7214 ± 0.039
	D&D	0.7931 ± 0.022	0.8376 ± 0.020	0.8248 ± 0.028	0.8402 ± 0.024
	PROTEINS	0.8137 ± 0.017	0.8210 ± 0.039	0.7834 ± 0.043	0.8349 ± 0.038
<i>k</i> -GNN	ENZYMES	0.5152 ± 0.111	0.5728 ± 0.063	0.5730 ± 0.077	0.5715 ± 0.080
	D&D	0.7562 ± 0.021	0.7785 ± 0.020	0.7948 ± 0.022	0.7864 ± 0.021
	PROTEINS	0.7636 ± 0.043	0.7518 ± 0.034	0.7814 ± 0.028	0.7857 ± 0.026