

# Multigrid Solver for Laplacian Linear Systems on Power-Law Graphs

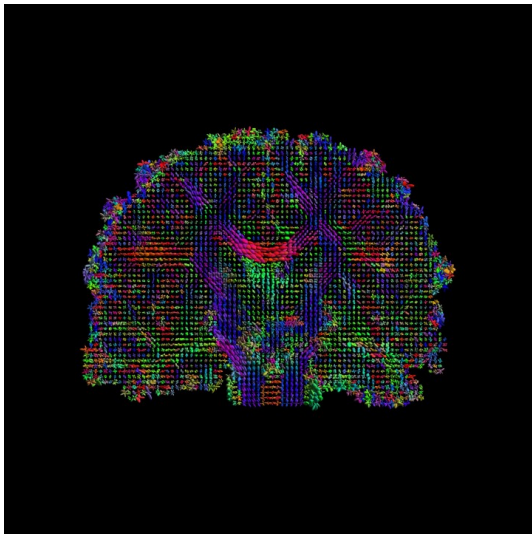
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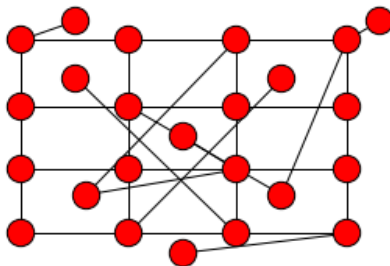




Human brain connectome; coronal view [5]

# Foundation

**Graph:**  $G = (N, E)$  where  $N$  is the set of nodes and  $E$  is the set of edges.

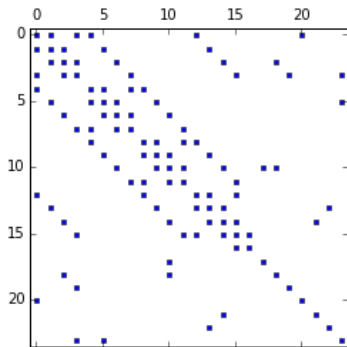


# Foundation

Laplacian matrix of  $G$ :

$$L = D - A$$

$D$ : diagonal matrix of node degrees,  $A$ : adjacency matrix of  $G$



# Why Do We Care About the Laplacian?

- Laplacian operator inverse is weighted average of paths through a network:

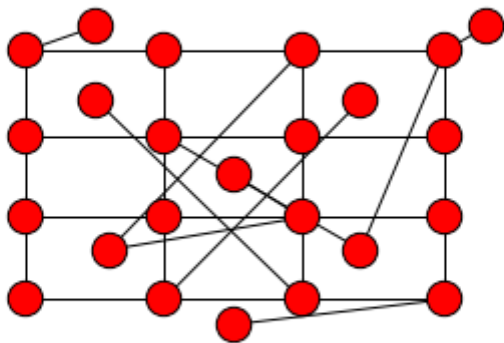
$$b + Lb + L^2b + L^3b + \dots = \sum_{i=0}^{\infty} L^i b = (I - L)^{-1} b$$

- PageRank linear system:

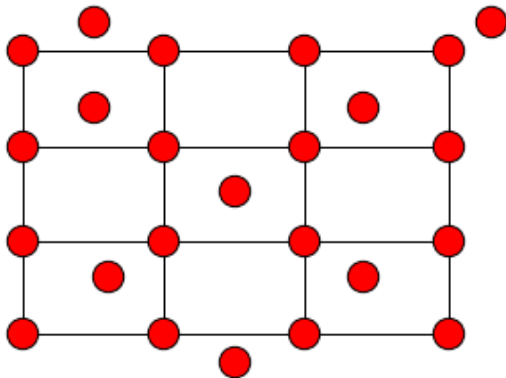
$$(I - \alpha L)x = (1 - \alpha)b$$

- Partition the graph into large planar subgraph and small teleportation subgraph
- Solve the large planar part with algebraic multigrid
- Solve the teleportation part directly
- Linear algebra to combine and solve the entire linear system

# Graph Partitioning



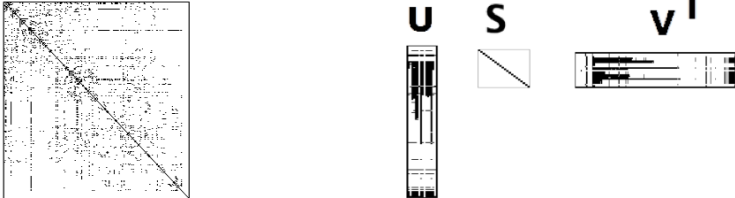
# Graph Partitioning



Max locally-connected subgraph using Chun-Lu [1]



# Linear System Solve

$$L = P_L + U S V^T$$


The diagram illustrates the LU decomposition of a matrix  $L$ . The matrix  $L$  is shown as a sparse matrix with a diagonal line. The permutation matrix  $P_L$  is shown as a small square matrix. The lower triangular matrix  $U$  is shown as a tall, narrow matrix with non-zero entries on the diagonal and below. The small square matrix  $S$  is shown as a small square matrix. The row vector  $V^T$  is shown as a horizontal row of non-zero entries.

# Linear System Solve

## Definition (Sherman-Woodberry-Morrison)

$$(P_L + USV^T)^{-1} = P_L^{-1} - P_L^{-1}U(S^{-1} + V^T P_L^{-1}U)^{-1}V^T P_L^{-1}$$

$$Lx = b$$

$$x = L^{-1}b$$

$$x = (P_L + T_L)^{-1}b$$

$$x = (P_L + USV)^{-1}b$$

$$x = (P_L^{-1} - P_L^{-1}U(S^{-1} + VP_L^{-1}U)^{-1}VP_L^{-1})b$$

$$x = P_L^{-1}b - P_L^{-1}U(S^{-1} + VP_L^{-1}U)^{-1}VP_L^{-1}b$$

# Current Solution Approaches

Multigrid on entire graph: LAMG, CMG, Cascadic

- How do you coarsen the grid? Based on heuristics

Brute force factorization

- Slow and memory inefficient

# Partitioning Complexity

- Graph partitioning complexity relies on the degree distribution of the graph
- Power law degree graphs with degree sequence:

$$P(d) \sim d^{-\gamma}$$

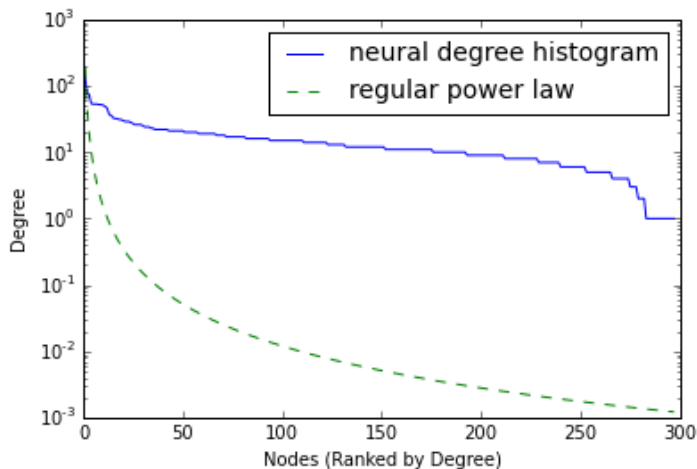
$$E[d] = \frac{\zeta(\gamma-1)}{\zeta(\gamma)}$$

- Partitioning is  $O(\text{iter.} \times \text{edges} \times E[d]^3)$

# Solve Complexity

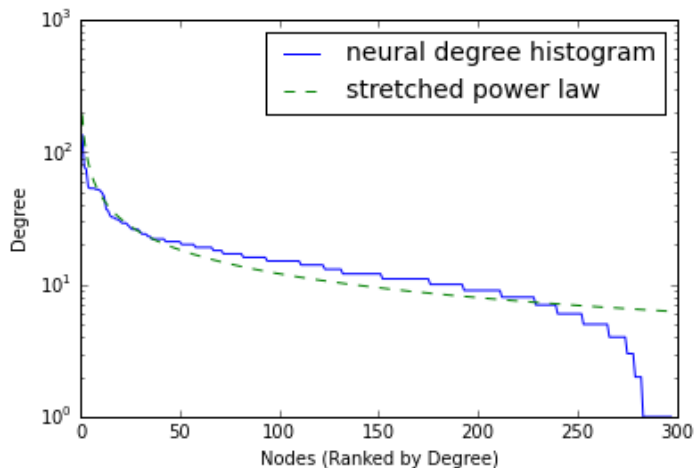
Operation	O(FLOPs)
$USV = T_L$	$O(n^3)$
$S^{-1}$	$O(r)$
$y = P_L^{-1}b$ (MG)	$O(n)$
$y_1 = Vy$	$O(rn)$
$Q = P_L^{-1}U$ ( $r$ MG solves)	$O(rn)$
$Q_1 = VQ$	$O(r^2n)$
$Q_2 = S^{-1} + Q_1$	$O(r^2)$
$y_2 = Q_2^{-1}y_1$	$O(r^3)$
$y_3 = Uy_2$	$O(rn)$
$y_4 = P_L^{-1}y_3$ (MG)	$O(n)$
$x = y - y_4$	$O(n)$

# Power Law vs. Neural Network Degree Sequence



$$\gamma = 2.1$$

# Stretched Power Law



stretched degree is 0.6

# Graphs and Results

Graph (nodes, edges)	Avg. Deg.	Rank $T_L$	Part. (s)	Iter.
Neural (297, 2148) [6, 7]	14.46	22	1.6	3
Metabolic (453, 2025) [2]	8.94	49	2.3	3
Protein (912, 22738) [4]	49.86	26	53	2
Facebook (4039, 88234) [3]	43.69	180	480	3
Power (4941, 6594) [6]	2.67	4284	.33	3



# Solve Component Results

These are the limiting solve components according to theoretical complexity and timed results:

Operation	$O(r, n)$	Neur.	Meta.	Prot.	FB	Pow.
$USV = T_L$	$O(n^3)$	.0334	.0737	.4183	38.47	72.86
$Q = P_L^{-1} U (r \times \text{MG})$	$O(rn)$	.3292	.7360	.5190	23.11	106.9
$Q_1 = VQ$	$O(r^2n)$	.0006	.0036	.0013	.4124	285.1
$y_2 = Q_2^{-1} y_1$	$O(r^3)$	.0023	.0025	.0013	.0099	86.49
<b>Total</b>	$O(n^3)$	<b>.51</b>	<b>.966</b>	<b>1.44</b>	<b>64.46</b>	<b>560</b>

# What Do These Solutions Mean?

## *C. Elegans* data:

- Important regions in the worm nervous system
- Key metabolic processes
- Unobserved protein-phenotype relationships

## Facebook friend networks:

- Origins of influence in a social group

## Power Grid:

- Flow of electricity through urban infrastructure

# Conclusions and Future Work

- Method to solve the graph laplacian linear system using partitioning and multigrid
- Optimize the solve operations and write faster, parallelizable code
- Find appropriate linear systems to solve the graphs I have
- Test larger graphs; more complex biological systems

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