Graph Bootstrap Percolation

S. Martine: T. Wang

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Graph Bootstrap Percolation

Case studies on selected graphs and algorithmic analysis

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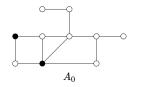
Empirica findings

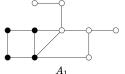
Result

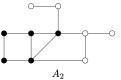
Consider the following activation process:

- Let G = (V, E) be a graph, whose vertices may be either in an "active" or "inactive" state. Fix a threshold $k \in \mathbb{N}$, and choose a seed set $A_0 \subseteq V$.
- Denoting the adjacent vertices, or neighbors of a given vertex v by N(v), activations (also called infections) spread through the system as follows:

$$A_{i+1} = A_i \cup \{v \in V : |N(v) \cap A_i| \ge k\}$$







A simple example

Motivation

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Bootstrap percolation can model spread in real life.

- Vertices and edges represent people and relationships, while the percolation process represents the spread of ideas, rumors, or trends.
- Which groups at minimum should be advertised to for maximal influence throughout a community?
- How can we reduce the spread of ideas or influence?

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In the language of bootstrap percolation:

- Minimum contagious sets
 - What is the smallest possible seed set A_0 (and its size $|A_0|$) required to eventually infect an entire graph?
- Spread minimization
 - If we must initially infect a fixed number of vertices (i.e., choose a seed set $A_0 \subseteq V$ of fixed cardinality), which vertices should be infected to minimize spread?

Minimum contagious sets

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Rocult

Let G(V, E) be a graph, and fix a threshold k for the percolation process.

- A contagious set is a seed set A_0 that eventually infects the entire graph (i.e., $\langle A_0 \rangle = V$).
- A contagious set is called minimum if there exists no smaller contagious set, and its size is denoted by m(G, k).
- Goal: Find m(G, k).
- One solution: First, compute all possible contagious sets by exhaustively finding all seed sets A_0 for which $\langle A_0 \rangle = V$. Then, m(G, k) is equivalent to the cardinality of the smallest A_0 (there may be more than one).

Spread minimization

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Let G(V, E) be a graph and fix a threshold k for the percolation process. If we must initially infect a fixed number of vertices (i.e., choose a seed set $A_0 \subseteq V$ of cardinality n), which vertices should be infected to minimize spread?

- For any seed set $A_0 \subseteq V$, we denote the set of all vertices that eventually become infected by $\langle A_0 \rangle$.
- Goal: Find a seed set A_0 of size n that minimizes $|\langle A_0 \rangle|$.
- One solution: Compute $\langle A_0 \rangle$ for all possible seed sets A_0 on the graph to find the smallest $|\langle A_0 \rangle|$ by exhaustion.

Challenges

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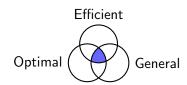
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- Spread minimization is believed to be NP-hard (though this remains unproven)
- This implies there is unlikely to be an efficient, optimal, general solution to the problem
- Hence, most approaches focus on specific cases (sacrificing generality) or finding a suboptimal solution
- Past work on the problem focuses on many different cases, supporting the suspected NP-hardness of the problem



Some interesting cases

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Results

In order to better understand bootstrap percolation, we worked through percolation processes on specific kinds of graphs.

- Complete graphs
- Petersen graph
- Bipartite graphs

Complete graphs

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Results

In K_n , the complete graph of n vertices, each vertex is adjacent to every other vertex. Given a seed set A_0 and threshold k, whether any other vertices can be infected is predetermined.

- If $|A_0| < k$, no vertices have k or more infected neighbors, so no spread occurs. Hence $|\langle A_0 \rangle| = |A_0|$.
- Otherwise, $|A_0| \ge k$, and all vertices have k or more infected neighbors. Therefore, the entire graph becomes infected in the next step, and $|\langle A_0 \rangle| = |V|$.



 K_7 , the complete graph of 7 vertices

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Result

The Petersen graph is a peculiar graph with 10 vertices and 15 edges. It is helpful to divide its vertices into two disjoint sets: the "inner" vertices and the "outer" vertices.



The Petersen graph

We examined outcomes for $|\langle A_0 \rangle|$ with threshold k=2 for different seed sets A_0 .

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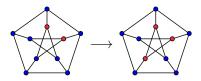
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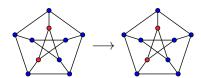
Reculto

Considering the "inner" vertices when $|A_0| = 2$:

Selecting two nearby vertices results in at most one subsequent infection.



■ Selecting not so nearby vertices as A_0 results in no further infections, and $|\langle A_0 \rangle| = |A_0|$.



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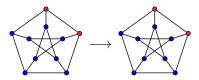
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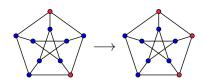
Reculto

Considering the "outer" vertices when $|A_0| = 2$:

Selecting two nearby vertices results in no further infections.



■ Selecting not so nearby vertices as A_0 results in at most one subsequent infection, and $|\langle A_0 \rangle| = |A_0|$.



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Why does this happen?

- There is an automorphism on the Petersen graph that preserves its shape, and maps the inner vertices to the outer vertices.
- Any choice of two nearby inner vertices maps to two outer vertices that are not so nearby.
- Therefore, the opposite rule applies after applying the automorphism.

Bipartite graphs

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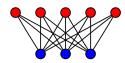
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A bipartite graph $K_{m,n}$ is a graph whose vertex set V can be split into two disjoint subsets A and B, where |A| = m and |B| = n.

- A bipartite graph is called complete if every vertex in A connects to every vertex in B.
- Percolation on the complete bipartite graph behaves similarly to the complete graph K_n .
- We explored percolation on bipartite graphs by running the percolation process on complete bipartite graphs with different combinations of *m* and *n*.



The complete bipartite graph $K_{3,5}$

Complete bipartite graphs

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Results

For $m, n \ge k$, a large enough and well distributed seed set A_0 will infect the entire graph $K_{m,n}$.

- If $|A_0| < k$, every vertex in $K_{m,n}$ has less than k infected neighbors and remains inactive.
- If $|A_0| \ge k$, A_0 can be entirely in subset A or B to infect the entire graph.
 - If A_0 is chosen to be in subset A, subset B will have $|A_0| \ge k$ infected neighbors and its vertices become activated during the first iteration of percolation.
 - During the second iteration, all vertices in A will now have $n \ge k$ infected neighbors and be activated.

General bipartite graphs

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For any general bipartite graph $K_{m,n}$, $m \ge n$, if $\lceil \frac{|A_0|}{2} \rceil < k$, then we can choose a seed set such that no new vertices are activated during the percolation process.

- For a graph where $m, n \ge \lceil \frac{|A_0|}{2} \rceil$, split the seed set exactly in half. If the seed set is odd sized, let the two halves differ by 1 vertex.
- For a graph where $n < \lceil \frac{|A_0|}{2} \rceil$, activate all of subset B and as much of subset A as necessary.

Erdős-Rényi random graphs

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The Erdős–Rényi graph is a random graph represented by G = G(n, p). n is the number of vertices and p is the associated probability.

- Edges of all possible combinations of vertices are added to
 G with probability p
- We examined only the cases in which this process creates a connected graph – any isolated vertex must be in the seed set to ever get activated
- Con: Does not accurately portray social communities

Example of Erdős-Rényi graphs

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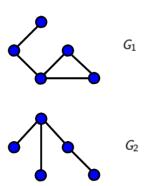
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- Two possible Erdős–Rényi graphs for G(5,0.3)
- All possible edges among 5 vertices are selected with probability 0.3



Stochastic block model

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Random graphs

A stochastic block model (SBM) graph is the union of two Erdős–Rényi graphs G_1 and G_2 . There is an additional probability q and two different communities G_1 and G_2 .

- We have a graph $G = (n_1, n_2, p_1, p_2, q)$
- \blacksquare n_i is the number of vertices per graph
- p_i is the probability per graph for intra-connected edges
- q is the probability for inter-connected edges between communities G_1 and G_2

Example stochastic block model graph

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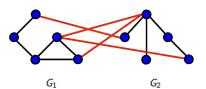
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Results

- We union the two Erdős–Rényi graphs from before
- Edges between vertices of different communities are added with probability q



Stochastic block model

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Results

In Bootstrap percolation on the stochastic block model (Torrisi, Giovanni Luca, Michele Garetto, and Emilio Leonardi., 2022), asymptotic bounds are given for p in terms of n and k.

$$1/n_i \ll p_i \ll 1/(n_i^{1/k})$$

Additionally, a critical number of seeds per community g_i is given in terms of p, n, and k.

$$g_i = \left(1 - \frac{1}{k}\right) \left(\frac{(k-1)!}{n_i p_i^k}\right)^{\frac{1}{k-1}}$$

■ When the the size of the seed set $|A_0| = g_i$, there is a "phase transition" in which the graph is rapidly infected

A brute-force approach

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We generated computations to observe the behavior of percolation in the SBM model which includes non-asymptotic cases. Generating these computations would also allow us to see how choosing a random seed set performs for percolation. We set the following parameters,

- $n_1 = n_2 \text{ and } p_1 = p_2$
- Sizes for *n* of 10, 50, 100, 500, 1000 vertices
- $2 \le |A_0| \le 4$ and $2 \le k \le 4$
- k is bounded below by $|A_0|$
- q is a constant multiple of p
- 40 p-values bounded between $1/n_i$ and $1/(n_i^{1/k})$

SBM computations, 10 vertices

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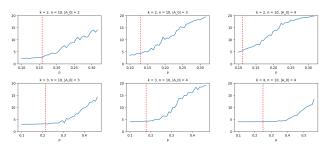
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Randomly chosen seed sets for n = 10

SBM computations, 1000 vertices

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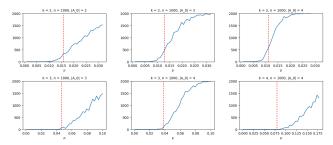
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Randomly chosen seed sets for n = 1000

SBM Results

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Result

The red dotted line indicates when the critical number of seeds g_i is reached and equal to $|A_0|$ for some value of p. This can be derived by solving for p in $g_i = \left(1 - \frac{1}{k}\right) \left(\frac{(k-1)!}{n_i p_i^k}\right)^{\frac{1}{k-1}}$. For a randomly chosen seed set, it can be seen that,

- When $|A_0| < g_i$, there is little to no spread
- When $|A_0| > g_i$, we see the phase transition in which $|\langle A_0 \rangle|$ gets larger
- So far, it looks like percolation in the SBM model performs accordingly to defined bounds!

An algorithmic approach

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An attempt to choose A_0 based on degree of vertices, in a few different ways. We hoped to minimize the known number of edges between active and inactive vertices

- There are two simple cases picking the largest degree vertices, and picking the smallest degree vertices
- Can either also choose to infect the neighbors of the chosen vertices or not
- Looked into our own algorithm that chooses the vertex with "greatest degree less than remaining size of seed set"

Algorithms, 10 vertices

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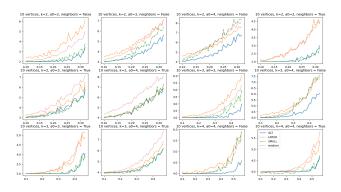
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The algorithms on random graphs of 10 vertices

Algorithms, 1000 vertices

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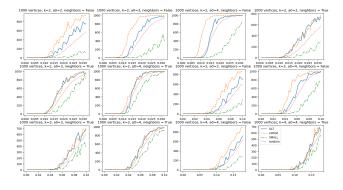
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The algorithms on random graphs of 1000 vertices

Results

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Empirically, choosing the vertices with smallest degree and not considering neighbors is the best algorithm for minimizing activation of a random graph

- Considered the same p-values as in the SBM model
- 10 vertex case is unique, potentially due to either
 - Higher variance
 - Both $|A_0|$ and k being close to the number of vertices

Next Steps

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We plan to continue looking into the algorithmic approach to better understand the probability behind why certain algorithms perform as they do, and how likely algorithms are to perform within a certain threshold.

Questions

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Thank you! Any questions?