

The K-clique Percolation Method
Clique Clustering

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Outline

Percolation

The Phase TransitionTM

Cliques

Clique Percolation Algorithm

Addenda



Section 1

Percolation

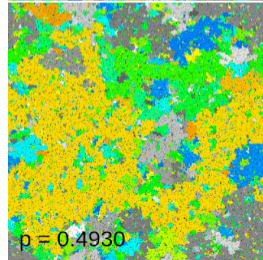
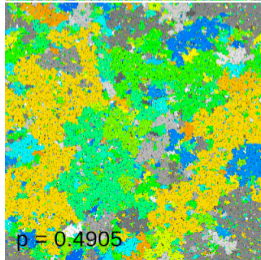
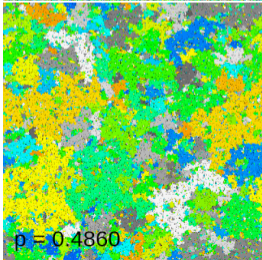
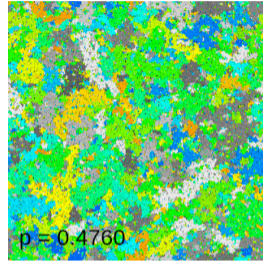
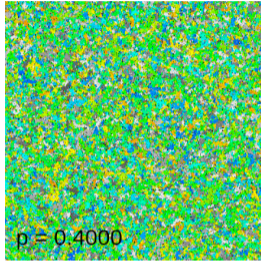
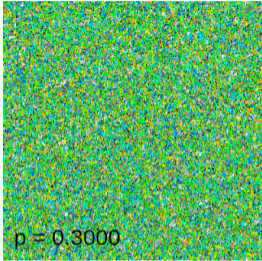


Percolating Water Problem

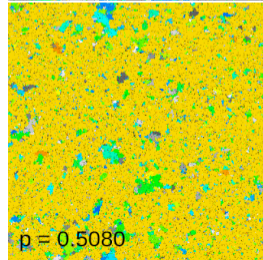
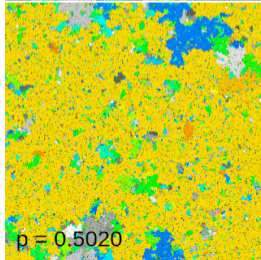
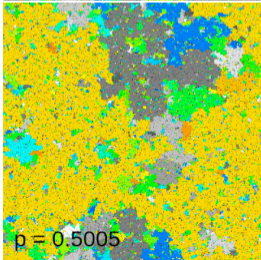
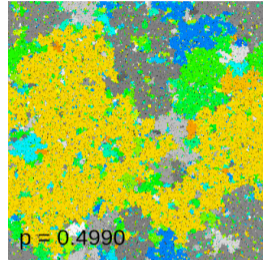
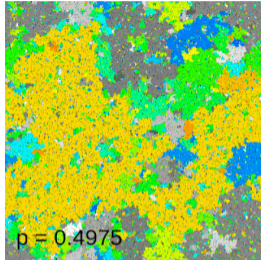
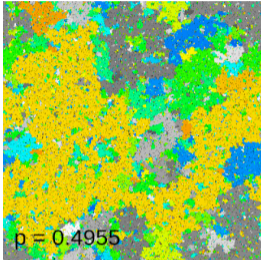
- Assume that some liquid is poured on top of some porous material. Will the liquid be able to make its way from hole to hole and reach the bottom?
 - ▶ **Model:** three-dimensional network of $n \times n \times n$ vertices, usually called "sites", in which the edge or "bonds" between each two neighbors may be open or closed
 - ▶ **Probabilistic:** A bond is open with probability p and closed $1 - p$.
- For a given p , what is the probability that an open path exists from the top to the bottom?



Below 50% Openness



Above 50% Openness



Section 2

The Phase Transition™



New Topic: Random Graphs [FK23]

- **Idea:** Define some parameters and generate a graph probabilistically
- **Erdős–Rényi Model:** The most common model
- $G(n, p)$: Add n nodes, then add edges with probability p
- $G(n, M)$: Add n nodes, then pick uniformly from all configurations of M edges

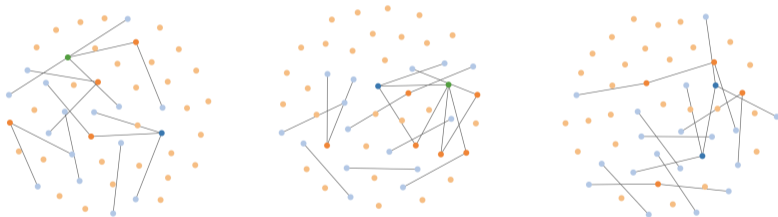


The "Percolated" Graph

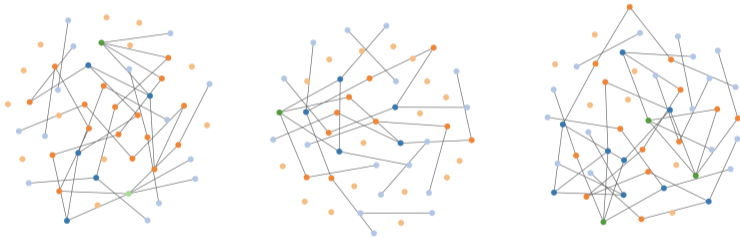
- The ER graph's definition is closely related to the percolation problem defined earlier
- **$G(n,p)$** : each edge has a fixed probability of being present or absent, independently of the other edges
- **An ER graph also has a percolation constant!**
- The evolution of the ER graph structure with increasing p can be very precisely proven [not in this presentation] but it is dominated by the same critical percolation constant of $p = \frac{1}{n}$



Erdős–Rényi Phase Transition: $p = 75\% \frac{1}{n}$



Erdős–Rényi Phase Transition $p = 125\% \frac{1}{n}$



Generalizing

- The giant connected component is made up of nodes connected to *each other*
- The giant connected component is made up of pairs of nodes connected to **other, possibly overlapping pairs of nodes**
- Can we find denser "clusters" by finding **cliques** connected to other cliques? **Yes!**

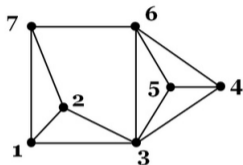


Section 3

Cliques



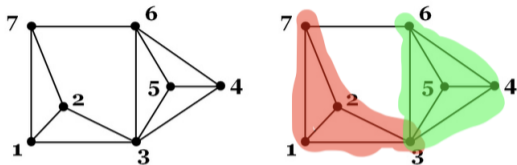
Cliques Review



- Some definitions for review:
- *Definition:* A clique c is a fully connected set of nodes, i.e. every pair of its vertices is connected by a link in the graph.
- *Definition:* A k -clique c_k is a clique of k nodes.
- *Definition:* Two k -cliques are said to be adjacent if and only if they share $k - 1$ nodes.



CPM Community



- *Definition:* A "clique percolation method" community is defined as the maximal union of k -cliques that can be reached from each other through a series of adjacent k -cliques
- How do we intuit what this means?
- Start with a clique "template", and "roll it over" on the graph. $\{1, 2, 7\}$ rolls over to $\{1, 2, 3\}$
- Alternatively, the "Group Chat of Theseus": take one node out of the clique, include a new one.
- Where are the 3-clique communities on this graph?



Questions?



Section 4

Clique Percolation Algorithm



Motivation

- We want to understand the structure of a graph - the "communities" within it
 - ▶ ex. social media - what topics or communities someone participates in or follows
 - ▶ ex. biology - identify related proteins within an interaction network
- Our approach (overlapping cliques) is an intuitive and deterministic definition of a community
- Allows for overlapping communities - many other methods don't do this
- Density requirement is freely adjustable via k



Algorithm Requirements

- *Subtask*: find the cliques
 - ▶ Either find all maximal cliques, or find all k -cliques in a graph
 - ▶ These are both NP-hard problems! Outside the scope of this lecture
 - ▶ A recent CPM paper uses a parallelized algorithm [Dan18]
- *Subtask*: storing the communities / connected components
 - ▶ We will use a **Union-Find data structure** (disjoint sets)
 - ▶ `UF.MakeSet()`: creates a new tree with one node p , corresponding to a new empty set, and returns p .
 - ▶ `UF.Find(p)` returns the root of the tree
 - ▶ `UF.Union(r_1, \dots, r_l)`: performs the union of trees represented by their roots r_i by making one root the parent of all others.



Pseudocode

BASIC CPM ALGORITHM(G):

```
1: UF  $\leftarrow$  Union-Find data structure
2: DICT  $\leftarrow$  []
3: for each  $k$ -clique  $c_k \in G$  do:
4:    $S \leftarrow \{\}$ 
5:   for each  $(k - 1)$ -clique  $c_{k-1} \subset c_k$  do:
6:     if  $c_{k-1} \in \text{DICT.KEYS}()$  then
7:        $p \leftarrow \text{UF.FIND}(\text{DICT}[c_{k-1}])$ 
8:     else
9:        $p \leftarrow \text{UF.MAKESET}()$ 
10:       $\text{DICT}[c_{k-1}] \leftarrow p$ 
11:       $S \leftarrow S \cup \{p\}$ 
12:    $\text{UF.UNION}(S)$ 
```



Questions?



Section 5

Addenda



Percolation Threshold

- There are also threshold thresholds for clique communities. What edge probability p is the threshold to producing one giant connected community in an ER graph?
- Intuition: going back to the "rolling-over" analogy. Within a "rolling", at the threshold, there should be, in expectation, one adjacent clique at each clique, to "continue" to build the cluster.
- $(k - 1) \times (N - k - 1)p_c^{k-1} = 1$: candidate vertices to remove \times candidate vertices to add
- For large N , this comes out to $N(k - 1)p_c^{k-1} = 1$ [Der05]

$$p_c(k) = \frac{1}{[N(k - 1)]^{\frac{1}{k-1}}} \quad (1)$$



Optimizations

- The given CPM algorithm is prohibitively expensive in memory because we store large number of $(k - 1)$ -cliques
- *Optimization:* instead of disjoint sets of $(k - 1)$ cliques, consider non-disjoint sets of z -cliques s.t. $z < (k - 1)$ [Bau22]
- why are there fewer z -cliques? Consider binomial theorem, $k \ll M$
- this leads to an *approximation* of the previous algorithm, but it is otherwise very similar and produces very similar results



Questions?



Bibliography



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