

Random Sums and Graphs

Probability

Events, probabilities,
conditional probabilities,
union bound

Random variables,
expected value,
linearity of expectation
Markov's inequality

Algorithms

Approx. 3-SAT,

Big idea: we only want the average

$$\text{Linearity of Expectation} \\ E[X+Y] = E[X] + E[Y] \quad (\text{independently useful}) \\ \text{"average sum = sum of averages"}$$

let $Y_i = \begin{cases} 1 & \text{if } i\text{th clause satisfied} \\ 0 & \text{otherwise} \end{cases} \quad (i=1, \dots, m)$

$$E[Y_i] = 7/8 \quad (1 \text{ clause case})$$

$$E[\text{\# clauses satisfied}] = E\left[\sum_{i=1}^m Y_i\right] = \sum_{i=1}^m E[Y_i] = 7/8 m$$

let $Y_i = \begin{cases} 1 & \text{if } i\text{th clause satisfied} \\ 0 & \text{otherwise} \end{cases}$

$$Z = \sum_{i=1}^m Y_i = \text{\# clauses satisfied}$$

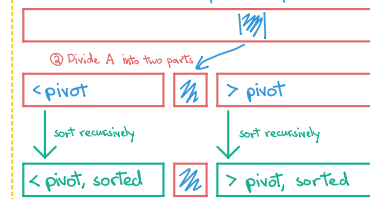
$$E[Z] = \frac{1}{2} E[Z | x_1 = \text{true}] + \frac{1}{2} E[Z | x_1 = \text{false}]$$

$\Rightarrow$ either $E[Z | x_1 = \text{true}]$ or $E[Z | x_1 = \text{false}]$ is $\geq E[Z]$!

QuickSort,

Random divide and conquer

① pick random 'pivot'



Expected running time of QuickSort

For each $i, j \in [n]$ w/ $i < j$,

$$X_{ij} = \begin{cases} 1 & \text{if } i\text{th smallest element is compared to } j\text{th smallest} \\ 0 & \text{otherwise} \end{cases}$$

$\sum_{i < j} X_{ij}$ = total # comparisons made by the algorithm
 $\approx$ total running time

$E[\sum_{i < j} X_{ij}]$ = average running time

$$E[\sum_{i < j} X_{ij}] = \sum_{i < j} E[X_{ij}]$$

now analyze $E[X_{ij}]$ in isolation

$$E[X_{ij}] = \Pr[X_{ij} = 1] \\ = \Pr[\text{\#th element is compared to } j\text{th}]$$



i th elem is compared w/ j th
 $\Leftrightarrow$ i th or j th elem selected before any element in between happens w/ probability $\frac{2}{j-i+1}$

$$E[X_{ij}] = \frac{2}{j-i+1}$$

$$E[X_{ij}] = \sum_{i < j} \frac{2}{j-i+1}$$

$$\leq 2n \left(\frac{1}{2} + \frac{1}{3} + \dots + \frac{1}{n} \right)$$

$$\leq O(n \log n) \quad \text{with harmonic series}$$

Quickselect (HW)

Probability

Hash Functions,

Universal Hash Functions

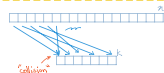
$h: [n] \rightarrow k$ s.t. for all pairs $i, j \in [n]$,
 $\Pr[h(i) = h(j)] \leq \frac{1}{k}$ (much weaker than ideal hash)

e.g. $h(x) = (ax + b) \bmod p \bmod k$

where • p prime, $> k$

• a in $\{1, \dots, p-1\}$ unif. random

• b in $\{0, \dots, p-1\}$ unif. random



Markov's Inequality

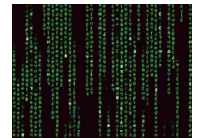
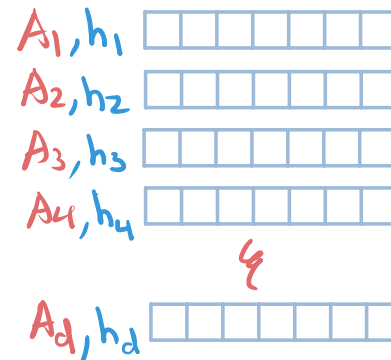
Markov's Inequality (special case)

Let $X \geq 0$ be nonnegative RV

$$\Pr[X \geq 2E[X]] \leq \frac{1}{2}$$

Algorithms

Frequency Estimation, Heavy Hitters (in streams)



estimate f_i w/
 $\min_{j=1, \dots, d} A_j[h_j(i)]$

Probability

k-wise independence,
higher order moments
(generalizing universal hash
functions and Markov's inequality)

Algorithms

Hash tables w/
chaining +
linear probing

k-wise independent hash functions

$$h: [U] \rightarrow [m]$$

s.t. for any k keys $x_1, x_2, \dots, x_k \in [U]$,
 k values $y_1, y_2, \dots, y_k \in [m]$,

$$P[h(x_1)=y_1, h(x_2)=y_2, \dots, h(x_k)=y_k] = \frac{1}{m^k}$$

eg. $h(x) = a_0 + a_1x + a_2x^2 + \dots + a_{k-1}x^{k-1} \pmod{p}$

where $p \geq U$ prime,

$a_0, a_1, \dots, a_{k-1} \in [p]$ indep, uniformly random

Important lemma

k-wise independent random variables

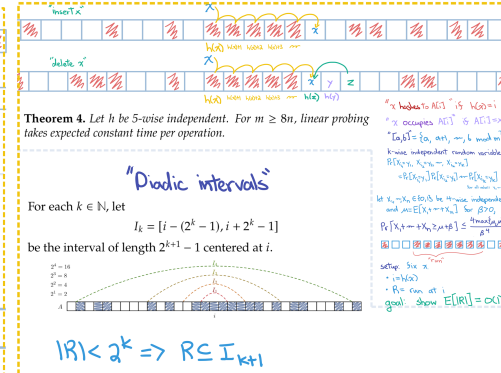
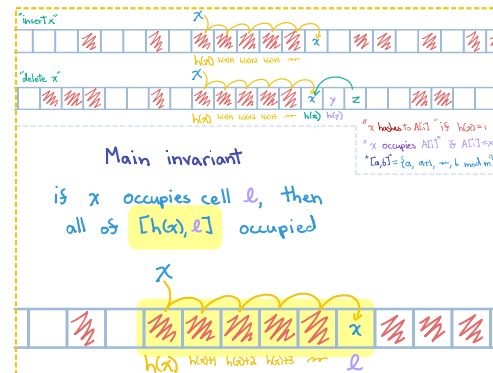
$$P[X_1=y_1, X_2=y_2, \dots, X_k=y_k]$$

$$= P[X_1=y_1] P[X_2=y_2] \dots P[X_k=y_k]$$

for all values $y_1, \dots, y_k$

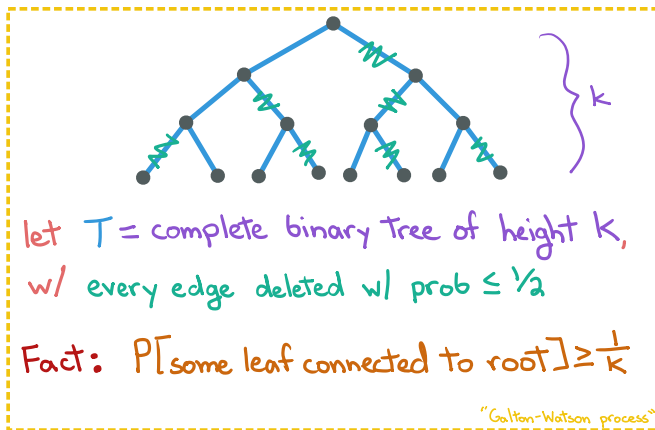
let $X_1, \dots, X_n \in \{0, 1\}$ be k -wise independent
and $\mu = E[X_1 + \dots + X_n]$. then for $\beta > 0$,

$$P[X_1 + \dots + X_n \geq \mu + \beta] \leq \frac{4 \max\{\mu, \mu^2\}}{\beta^4}$$



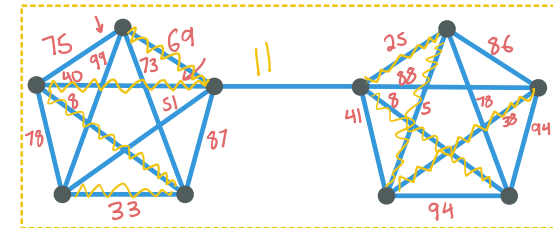
Probability

Galton-Watson processes



Algorithms

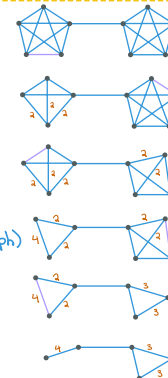
Random Contractions for min cut



Randomized contractions

Input: $G=(V,E)$, weights $c \in \mathbb{R}_{>0}^E$
(unweighted)

- ① while $|E| > 1$
 - Ⓐ sample $e \in E$ in proportion to c
 - Ⓑ "contract" e
- ② return set of edges (in original graph) contracted to remaining edge

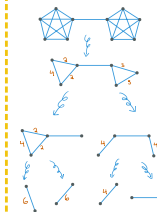


Amplification by branching

Instead of restarting at the beginning, restart partway through

High-level algorithm

- ① randomly contract until prob. of ruining the min-cut is $\sim 1/2$
- ② Branch:
make two copies of current graph
recurse on both copies



PLAYER	Red	1
BRIDGE	POWDER	10
THE	104	

YES NO

Would you like to DRIVE the game?

Probability

Multiplicative
Chernoff Bound

Algorithms (/ results)

Random graphs
& the giant component

Probability

Multiplicative
Chernoff Bound

Algorithms (/ results)

Random graphs
& the giant component



Flip 100 coins.

50 come up heads.

Should you be surprised?

(Maybe a little: $\approx 8\%$ chance)



Flip 100 coins.

0 come up heads.

Should you be surprised?

Absolutely: $\frac{1}{2^{100}}$ chance



Flip 100 coins.

≤ 25 come up heads.

Should you be surprised?

Yes: $\approx 2.818 \times 10^{-7} \%$



Flip 10 coins.

≤ 4 come up heads.

Should you be surprised?

No: $\approx 37.7\%$



Flip 1000 coins.

$\leq 40\%$ come up heads.

Should you be surprised?

Yes: $\approx 1.364 \times 10^{-10}$ probability



10 coins.

≤ 4 heads.

$\approx 37.7\%$

1000 coins.

$\leq 40\%$ heads.

1.364×10^{-10} prob.

scale matters

Concentration phenomena

let $X_1, \dots, X_n \in [0, 1]$ be independent

$$\mu = E[X_1 + \dots + X_n]$$

Q: probability that

$X_1 + \dots + X_n$ is close to μ ?
(multiplicatively)

A: error probability exponentially small in n

Multiplicative Chernoff bound

let $X_1, \dots, X_n \in [0, 1]$ be independent

$$\mu = E[X_1 + \dots + X_n]$$

$$\epsilon \in (0, 1)$$

$$\bullet \quad P[X_1 + \dots + X_n \geq (1 + \epsilon)\mu] \leq e^{-\epsilon^2 \mu / 3}$$

$$\bullet \quad P[X_1 + \dots + X_n \leq (1 - \epsilon)\mu] \leq e^{-\epsilon^2 \mu / 2}$$

(proof at end if time permits)

Lecture 4 (last Thursday)

Probability

Multiplicative
Chernoff Bound

Algorithms (/ results)

Random graphs
& the giant component

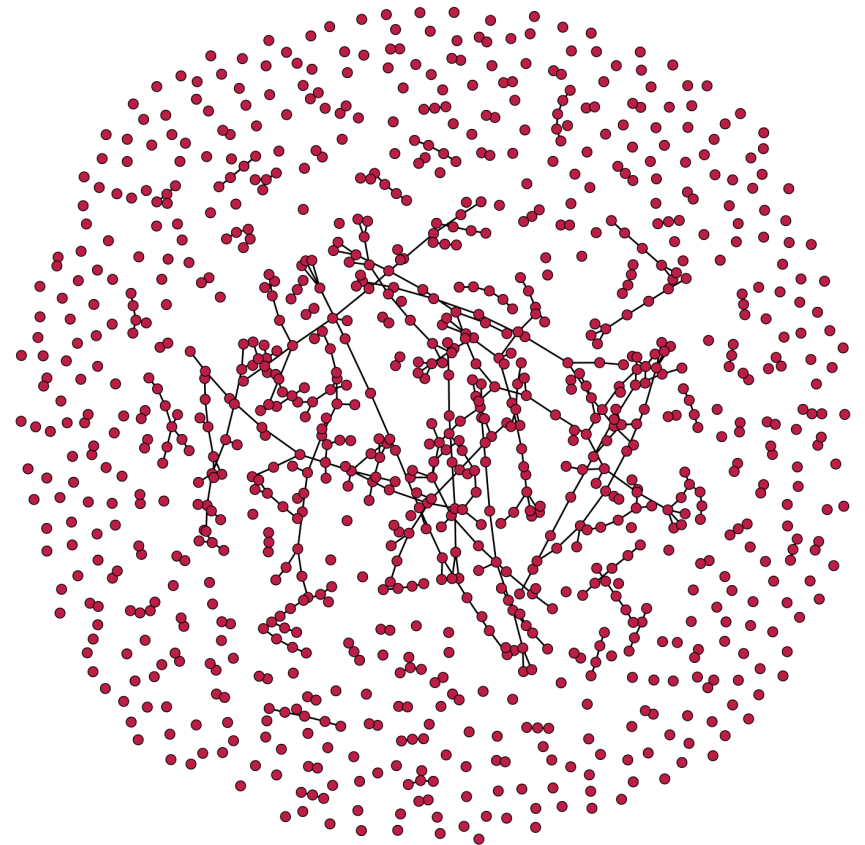
Erdős-Rényi graphs $G(n, p)$

(1959)

- n vertices
- each edge sampled independently
w/ probability p

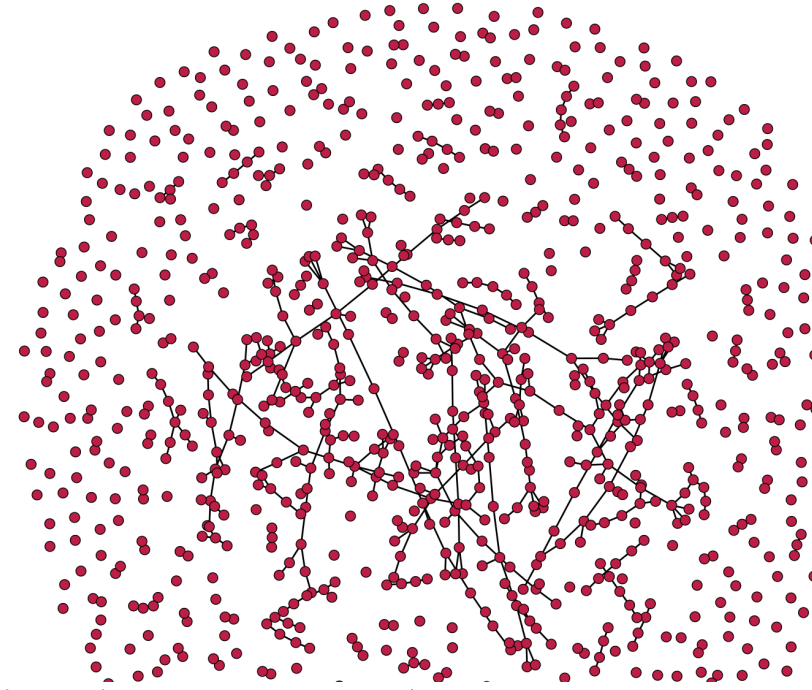
Q: what does such a graph
look like?

does it consistently have
certain properties?



Erdős-Rényi graphs $G(n, p)$

- n vertices
- each edge sampled independently w/ probability p



Theorem 5.3. Consider a random graph $G \sim \mathcal{G}(n, p)$ for $p = c/n$, where c is a constant.

1. If $c > 1$, then with high probability, there is exactly one connected component of G with $\Omega(n)$ vertices, and all other components have size $\leq O(\log n)$.
2. For $c < 1$, then with high probability, all connected components of G have size $< O(\log n)$.

"threshold phenomena"

Theorem 5.3. Consider a random graph $G \sim \mathcal{G}(n, p)$ for $p = c/n$, where c is a constant.

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Proof outline ($c > 1$)

- ① Gap theorem let $p = (1 + \epsilon)/n$. w/ high probability, all components have size
"small component" $\leq O\left(\frac{\log n}{\epsilon^2}\right)$ or "large component" $\geq \Omega(\epsilon n)$
- ② Existence of large component
- ③ Uniqueness of large component

① Gap theorem let $p = (1 \pm \epsilon)/n$. w/ high probability,
all components have size

$$\leq O\left(\frac{\log n}{\epsilon^2}\right) \quad \text{or} \quad \geq \Omega(\epsilon n)$$

Setup

fix vertex v .

let $C(v) \subseteq V$ = connected component of v



to analyze $C(v)$, imagine
revealing $C(v)$ w/ a search algo

Fix vertex v . $C(v) \subseteq V =$ connected component of v

"Search algo"

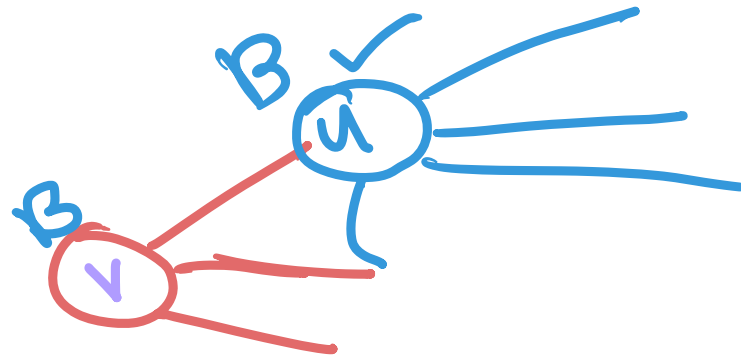
$A = \{\text{vertices known to be in } C(v)\}$ (initially $A = \{v\}$)

$B = \{\text{explored vertices}\}$ (initially $B = \emptyset$)

① while there is a vertex $v_i \in A \setminus B$

 ① add all neighbors of v_i to A

 ② add v_i to B



Indexed version of "Search algo"

$A_i = \{\text{vertices known to be in } C(v) \text{ after } i \text{ iter.}\}$
($A_0 = \{v\}$)

$B_i = \{\text{explored vertices after } i \text{ iterations}\}$ ($B_0 = \emptyset$)

1. For $i = 1, 2, \dots$ until $A_i = B_i$

a. Let $v_i \in A_{i-1} \setminus B_{i-1}$

b. $A_i = A_{i-1} \cup N(v_i)$

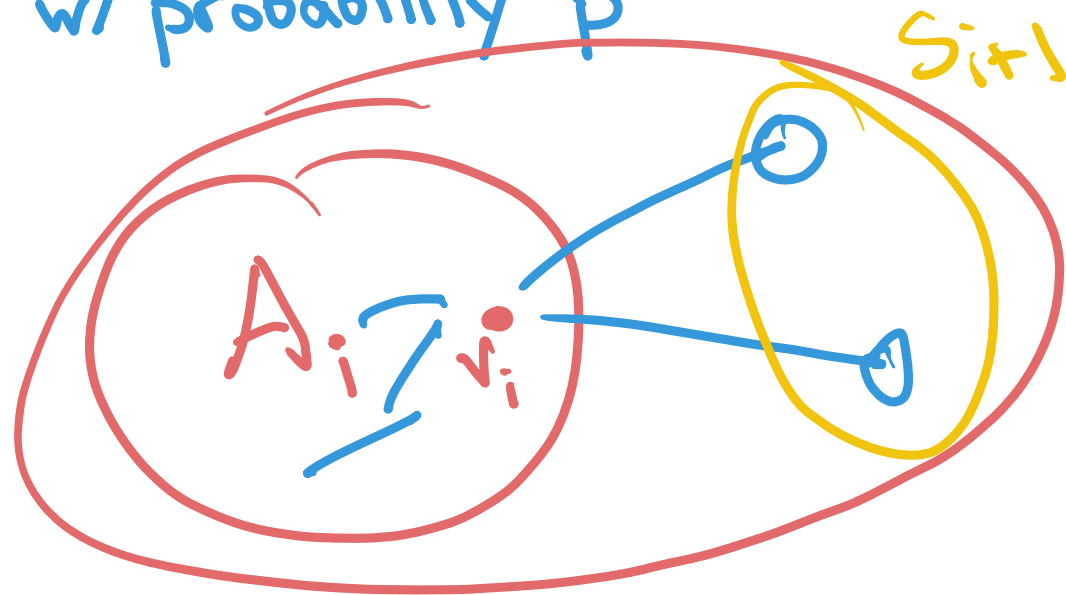
c. $B_i = B_{i-1} \cup \{v_i\}$

Observation 1

algo terminates iff $|A_i| = i$

Observation 2 $A_{i+1} = A_i \cup S_{i+1}$

where S_{i+1} samples each $v \in V \setminus A_i$ independently
w/ probability p



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c. $B_i = B_{i-1} \cup \{v_i\}$

$$p_i = \frac{(1+\epsilon)}{n} \cdot \frac{\epsilon n}{4} \leq \frac{\epsilon + \epsilon^2}{4}$$

(equivalent)

1. For $i=1,2,\dots$ ~~until $A_i = B_i$~~

a. Let S_i randomly sample $V \setminus A_{i-1}$ w/ probability p

b. $A_i = A_{i-1} \cup S_i$

let $i \leq \frac{\epsilon n}{4}$. $p = \frac{1+\epsilon}{n}$, $\epsilon > 0$

① Gap theorem let $p = (1+\epsilon)/n$ w/ high probability, all components have size $\leq O(\frac{\log n}{\epsilon^2})$ or $\geq \Omega(\epsilon n)$

Lemma 1. $E[|A_i|] \geq (1+\frac{\epsilon}{4})i$

$|A_0| = 1$

$E[|A_1|] = 1 + (n-1)p$

$E[|A_2|] =$

$1 + (1 - (1-p)^2)(n-1)$

$E[|A_i|] = 1 + \dots (1 - (1-p)^i)(n-1) \geq (1 + \frac{\epsilon}{4})i$

$(1-p)^i \leq e^{-pi} \leq 1 - pi + (pi)^2 \leq 1 - pi + \frac{(\epsilon+\epsilon^2)}{4} p_i$
 $e^x = 1 + x + \frac{x^2}{2!} + \frac{x^3}{3!} + \dots$

Multiplicative Chernoff bound
 let $X_1, \dots, X_n \in [0,1]$ be independent

$\mu = E[X_1 + \dots + X_n]$ $\epsilon \in (0,1)$

$P[X_1 + \dots + X_n \leq (1-\epsilon)\mu] \leq e^{-\epsilon^2 \mu / 2}$

1. For $i=1,2,\dots$ ~~until~~

a. Let S_i randomly sample

$V \setminus A_{i-1}$ w/ probability p

b. $A_i = A_{i-1} \cup S_i$

$$\text{let } i \leq \frac{\epsilon n}{4} \leq 1 - \frac{\epsilon}{4} p_i$$

① Gap theorem let $p = (1+\epsilon)/n$. w/ high probability, all components have size $\leq O(\frac{\log n}{\epsilon^2})$ or $\geq \Omega(\epsilon n)$

Lemma 1. $E[|A_i|] \geq (1+\frac{\epsilon}{4})i$

Lemma 2.

$$P[|A_i| \leq i] \leq e^{-\epsilon^2 i / 8}$$

$$X_j = \begin{cases} 1 & \text{if } j\text{th vertex in } A_i \\ 0 & \text{otherwise} \end{cases}$$

$$|A_i| = \sum X_j, \quad X_j \in [0, 1], \quad \text{indep.}$$

$$Pr[|A_i| \leq \frac{1}{1+\frac{\epsilon}{4}} E[|A_i|]] \leq Pr[|A_i| \leq (1-\frac{\epsilon}{8})\mu] \leq e^{-\frac{\epsilon^2}{64 \cdot 2} \mu} = e^{-(\epsilon^2/128)\mu}$$

Multiplicative Chernoff bound
let $X_1, \dots, X_n \in [0, 1]$ be independent

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$$P[X_1 + \dots + X_n \leq (1-\epsilon)\mu] \leq e^{-\epsilon^2 \mu / 2}$$

1. For $i=1, 2, \dots$ ~~until $|A_i| \geq i$~~

a. Let S_i randomly sample $V \setminus A_{i-1}$ w/ probability p

$$b. A_i = A_{i-1} \cup S_i$$

$$\frac{1}{1-x} \geq 1 + \frac{x}{2}$$

let $i \leq \frac{\epsilon n}{4}$.

Lemma 1. $E[|A_i|] \geq (1+\epsilon)i$

Lemma 2.

$$P[|A_i| \leq i] \leq e^{-\epsilon^2 i / 8}$$

for $i \geq 32 \log n / \epsilon^2$, $\uparrow \leq \frac{1}{n^4}$

by union bound over $\frac{32 \log n}{\epsilon^2} \leq i \leq \frac{\epsilon n}{4}$,

$$P\left[\frac{32 \log n}{\epsilon^2} \leq |A_i| \leq \frac{\epsilon n}{4}\right] \leq \frac{1}{n^3}$$

① Gap theorem let $p = (1+\epsilon)/n$. w/ high probability, all components have size $\leq O(\frac{\log n}{\epsilon^2})$ or $\geq \Omega(\epsilon n)$

Multiplicative Chernoff bound
let $X_1, \dots, X_n \in [0, 1]$ be independent

$$\mu = E[X_1 + \dots + X_n] \in (0, 1)$$

$$P[X_1 + \dots + X_n \leq (1-\epsilon)\mu] \leq e^{-\epsilon^2 \mu / 2}$$

1. For $i=1, 2, \dots$ until $|A_i|=i$
 - a. Let S_i randomly sample $V \setminus A_{i-1}$ w/ probability p
 - b. $A_i = A_{i-1} \cup S_i$

Theorem 5.3. Consider a random graph $G \sim \mathcal{G}(n, p)$ for $p = c/n$, where c is a constant.

Proof outline ($c > 1$)

1. If $c > 1$, then with high probability, there is exactly one connected component of G with $\Omega(n)$ vertices, and all other components have size $\leq O(\log n)$.
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✓ ① Gap theorem let $p = (1 + \epsilon)/n$. w/ high probability,
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 $\leq O\left(\frac{\log n}{\epsilon^2}\right)$ or $\geq \Omega(\epsilon n)$

② Existence of large component

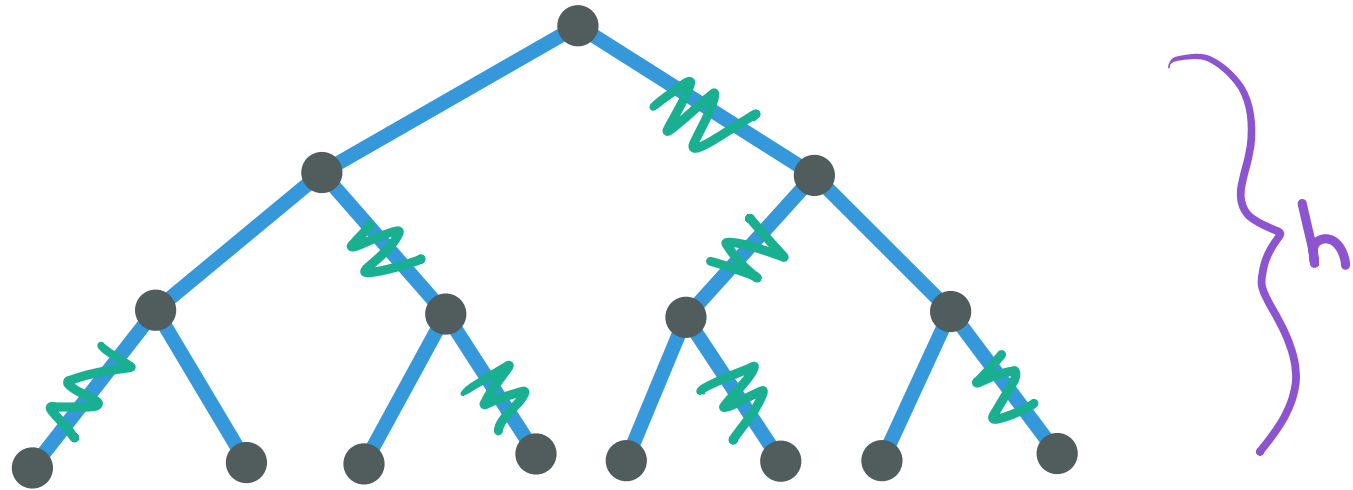
③ Uniqueness of large component

② Existence of large component

Lemma 5.5. *Let $p = (1 + \epsilon)/n$ for $\epsilon > 0$. Let $v \in V$ be a vertex. For all $3 \leq h \leq \epsilon n$, with probability at least $1/h$, v has at least $1 + h$ vertices in its connected component.*

Lemma 5.5. Let $p = (1 + \epsilon)/n$ for $\epsilon > 0$. Let $v \in V$ be a vertex. For all $3 \leq h \leq \epsilon n$, with probability at least $1/h$, v has at least $1 + h$ vertices in its connected component.

Last time

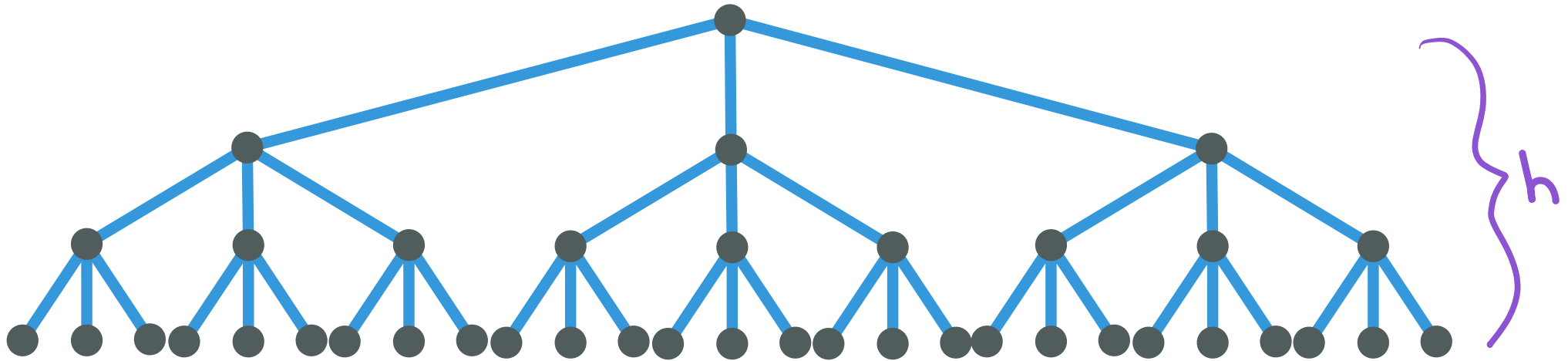


let T = complete binary tree of height h ,
w/ every edge deleted w/ prob $\leq 1/2$

Fact: $P[\text{some leaf connected to root}] \geq \frac{1}{h}$

"Galton-Watson process"

Lemma 5.5. Let $p = (1 + \epsilon)/n$ for $\epsilon > 0$. Let $v \in V$ be a vertex. For all $3 \leq h \leq \epsilon n$, with probability at least $1/h$, v has at least $1 + h$ vertices in its connected component.

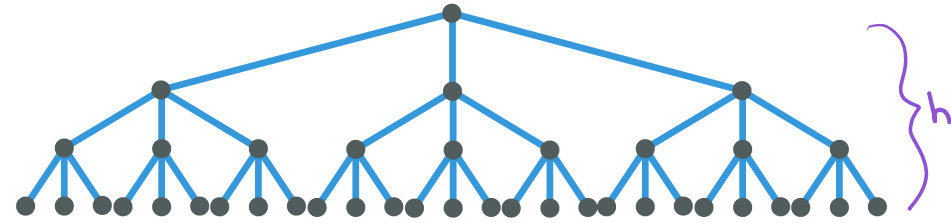


let $T =$ complete k -ary tree of height h ,
w/ every edge deleted w/ prob $\leq 1/k$

Fact: $P[\text{some leaf connected to root}] \geq \frac{1}{h}$

"Galton-Watson process"

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"Galton-Watson process"

Equivalent process

start w/ population of size 1

Each generation, each individual flips k coins
w/ prob. $1/k$ of heads

total # heads gives size of next generation

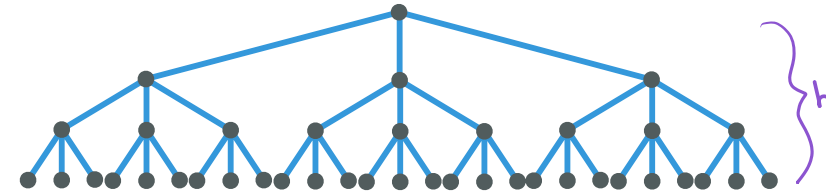
Fact: $P[\text{surviving } h \text{ generations}] \geq \frac{1}{h}$

Lemma 5.5. Let $p = (1 + \epsilon)/n$ for $\epsilon > 0$. Let $v \in V$ be a vertex. For all $3 \leq h \leq \epsilon n$, with probability at least $1/h$, v has at least $1 + h$ vertices in its connected component.

Recall search algo.

conditional on $|A_{i-1}| \leq h$,

A_i/A_{i-1} is like sampling
 $n-h \geq \frac{n}{1+\epsilon}$ children each
w/ prob. $p = \frac{1+\epsilon}{n}$



let T = complete k -ary tree of height h ,
w/ every edge deleted w/ prob $\leq 1/k$

Fact: $P[\text{some leaf connected to root}] \geq \frac{1}{h}$
"Galton-Watson process"

1. For $i=1, 2, \dots$ until $A_i = B_i$
 - a. Let $v_i \in A_{i-1} \setminus B_{i-1}$
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① Gap theorem let $p = (1 + \epsilon)/n$. w/ high probability,
all components have size
 $\leq O\left(\frac{\log n}{\epsilon^2}\right)$ or $\geq \Omega(\epsilon n)$

② Existence of large component

③ Uniqueness of large component

Imagine sampling $G \sim (n, p)$ as:

- ① sample $G_1 \sim (n/2, p)$
- ② sample $G_2 \sim (n/2, p)$
- ③ combine G_1 and G_2
- ④ sample cross edges



Multiplicative Chernoff bound

let $X_1, \dots, X_n \in [0, 1]$ be independent

$$\mu = E[X_1 + \dots + X_n]$$

$$\epsilon \in (0, 1)$$

$$\bullet \quad P[X_1 + \dots + X_n \geq (1 + \epsilon)\mu] \leq e^{-\epsilon^2 \mu / 3}$$

$$\bullet \quad P[X_1 + \dots + X_n \leq (1 - \epsilon)\mu] \leq e^{-\epsilon^2 \mu / 2}$$

(proof at end if time permits)

models, these graphs – however random – tend to be extremely consistent about certain properties.

Today we will study the $\mathcal{G}(n, p)$ random graph, sometimes called Erdős-Rényi graphs based on work by Erdős and Rényi [ER59; ER60]. A random graph from $\mathcal{G}(n, p)$ is an undirected graph over n vertices, where every edge is sampled independently with probability p . By now there is a large catalog of nontrivial and useful properties that, depending on p , are almost certain to appear or not appear in such a graph (for sufficiently large n). Moreover, Erdős and Rényi showed that these properties can vary dramatically with very small changes in p . Consider the following theorem.

Theorem 2.1. *Consider a random graph $G \sim \mathcal{G}(n, p)$ for $p = c/n$, where c is a constant.*

1. *If $c > 1$, then with high probability, there is exactly one connected component of G with $\Omega(n)$ vertices, and all other components have size $\leq O(\log n)$.*
2. *For $c < 1$, then with high probability, all connected components of G have size $< O(\log n)$.*

The parameter c above models the average degree (in expectation). The drama lies in the fact that a tiny change in the average degree c – from .999 to 1.0001 – flips the qualitative nature of a typical random graph from one of many tiny components to essentially one giant component. This is an example of a *threshold phenomena*; alternatively, a *nonlinear dynamic*. Such phenomena is not rare: it occurs in many situations in physics, as well as in models for epidemiology and social networks. Let us briefly mention — without claiming to be very precise — that the sensitivity to c gives some motivation for controlling the “reproductive number” when analyzing and preventing the spread of infectious diseases. The reproductive number is the expected number of healthy individuals that a sick individual effects.

We note that further research has obtained a much more refined and detailed understanding than stated in theorem 2.1. We refer the reader to [Bol98, Chapter 7] for further details and other results in this area.

We present the proof for $c > 1$. The second case of $c < 1$ is left to the reader as exercise 1.

2.1 Overview of the proof for $c > 1$

We will prove part 1 of theorem 2.1 in roughly three parts.

Part 1: the gap theorem. Observe that in theorem 2.1 above, regardless of the value of p , there are simply no “medium”-size components, such as a component of size $\sqrt{n}$ or of size $n/\log(n)$. The intermediate sizes are ruled out by the following “gap theorem”.

Lemma 2.2. *There is a universal constant $C > 0$, such that for all $\epsilon \in (0, 1)$, and for all $n > 0$ sufficiently large, and $p = (1 + \epsilon)/n$, we have the following. For a random graph*

designed elaborate randomized constructions of graphs and showed that with nonzero probability, they can possess certain counterintuitive, seemingly impossible properties. This general approach is now called *Ramsey theory*. Second, he showed that for natural random graph models, these graphs – however random – tend to be extremely consistent about certain properties.

Today we will study the $\mathcal{G}(n, p)$ random graph, sometimes called Erdős-Rényi graphs based on work by Erdős and Rényi [ER59; ER60]. A random graph from $\mathcal{G}(n, p)$ is an undirected graph over n vertices, where every edge is sampled independently with probability p . By now there is a large catalog of nontrivial and useful properties that, depending on p , are almost certain to appear or not appear in such a graph (for sufficiently large n). Moreover, Erdős and Rényi showed that these properties can vary dramatically with very small changes in p . Consider the following theorem.

Theorem 5.3. *Consider a random graph $G \sim \mathcal{G}(n, p)$ for $p = c/n$, where c is a constant.*

1. *If $c > 1$, then with high probability, there is exactly one connected component of G with $\Omega(n)$ vertices, and all other components have size $\leq O(\log n)$.*
2. *For $c < 1$, then with high probability, all connected components of G have size $< O(\log n)$.*

The parameter c above models the average degree (in expectation). The drama lies in the fact that a tiny change in the average degree c – from .999 to 1.0001 – flips the qualitative nature of a typical random graph from one of many tiny components to essentially one giant component. This is an example of a *threshold phenomena*; alternatively, a *nonlinear dynamic*. Such phenomena is not rare: it occurs in many situations in physics, as well as in models for epidemiology and social networks. Let us briefly mention — without claiming to be very precise — that the sensitivity to c gives some motivation for controlling the “reproductive number” when analyzing and preventing the spread of infectious diseases. The reproductive number is the expected number of healthy individuals that a sick individual effects.

We note that further research has obtained a much more refined and detailed understanding than stated in theorem 5.3. We refer the reader to [Bol98, Chapter 7] for further details and other results in this area.

We present the proof for $c > 1$. The second case of $c < 1$ is left to the reader as exercise 5.1.

5.2.1 Overview of the proof for $c > 1$

We will prove part 1 of theorem 5.3 in roughly three parts.