## Lecture 3: Block models \& information-computation gaps

In this lecture we give another simple application of the sum-of-squares paradigm, to the problem of recovering communities in stochastic block models with average degrees $\Omega(\log n)$. Sparse stochastic block models exhibit a notorious information-computation gap, and we describe the gap as well as give a heuristic derivation of the threshold. Some bibliographic remarks will be deferred to the end.

These notes have not been reviewed with the same scrutiny applied to formal publications. There may be errors.

## 1 Stochastic block models

Definition 1.1. The stochastic block model is a family of distributions over graphs, parametrized by integers $n$ (the number of vertices) and $k$ (the number of communities), and probabilities $p_{\text {in }}$ (internal edge probability) and $p_{\text {out }}$ (external edge probability).
One samples $G \sim \mathcal{G}\left(n, k, p_{\text {in }}, p_{\text {out }}\right)$ as follows:

1. Identify the vertex set $V(G)$ with [ $n$ ].
2. Sample a uniform partition of $[n]$ into $k$ sets or "communities" of equal size. Let $X \in\{0,1\}^{n \times k}$ represent this partition, with $X_{i c}=1$ [ vertex $i \in$ community $\left.c\right]$.
3. For each pair $i \neq j \in[n]$ independently, add edge $(i, j)$ to $E(G)$ with probability $p_{\text {in }}$ if $i, j$ are in the same community, and with probability $p_{\text {out }}$ otherwise.

In this lecture, we will focus on the case when $k=\Theta(1)$ and $\min \left(p_{\text {in }} n, p_{\text {out }} n\right)=\Omega(1)$. Some of what we prove can be extended to other regimes too.

These simple models are employed in multiple contexts; they are basic models for social networks, in which each partition can model a community of people (people are more likely to know other people from the same community). $\mathcal{C}\left(n, k, p_{\mathrm{in}}, p_{\text {out }}\right)$ is a primitive model for a social network, to be sure, but network science is hard, and if you want to evaluate a clustering algorithm (or some other sort of algorithm), it's not a bad place to start. Block models can also model physical systems, in which we think of each partition as a type of particle, and particles of the same type are either attracting ( $p_{\text {in }}>p_{\text {out }}$ ) or repulsive ( $p_{\text {in }}<p_{\text {out }}$ ).

Remark 1.2. Block models are sometimes defined more flexibly; the model above is sometimes referred to as the "symmetric block model." In the most general case, we allow communities of different sizes, and distinct probabilities between each pair of communities, $\left\{p_{a, b}\right\}_{a, b \in[k]}$.

Even in the symmetric block model, sometimes the partition $X$ is chosen so that the community of each $i \in[n]$ is chosen from $\operatorname{Unif}([k])$ independently, so that the parts in the partition are not even. This is sometimes more convenient for calculations, and when $k=\Theta(1)$, the differences between the models are usually qualitatively negligible.

Algorithmic questions on block models. There are two algorithmic questions typically associated with block models:

1. Recovery: given $G \sim \mathcal{C}\left(n, k, p_{\text {in }}, p_{\text {out }}\right)$, recover $X$ (either exactly or approximately).
2. Detection: you are given a graph $G$, and you wish to hypothesis test between the two hypotheses: $G \sim \mathcal{H}_{0}=\mathcal{C}\left(n, \frac{1}{k} p_{\text {in }}+\frac{k-1}{k} p_{\text {out }}\right)$, or $G \sim \mathcal{H}_{1}=\mathcal{G}\left(n, k, p_{\text {in }}, p_{\text {out }}\right)$. In the null hypothesis $\mathcal{H}_{0}$, there is no community structure.
In what follows, we will give an SoS algorithm for recovery in the regime $p_{\text {in }}, p_{\text {out }}=\Omega\left(\frac{\log n}{n}\right)$, and we will discuss a notorious information-computation gap for detection in the regime $p_{\text {in }}, p_{\text {out }}=O\left(\frac{1}{n}\right)$.

## 2 Sum-of-squares paradigm for recovery in block models

In the recovery problem, we wish to recover the partition $X$ which maximizes $\operatorname{Pr}[X \mid G]$, the maximum likelihood estimator. Here, the maximum likelihood estimate is not convenient to work with, and in particular the constraint that $X$ is the maximum likelihood estimator cannot be easily expressed as a low-degree polynomial constraint. Instead, we'll consider the maximum a-posteriori estimator, or the MAP estimator,

$$
Y^{*}=\underset{Y \text { balanced } k \text {-partition }}{\arg \max } \operatorname{Pr}[G \mid Y]
$$

We'll prove that if $G$ is sampled from partition $X$, then with high probability $\left(Y^{*}\right)\left(Y^{*}\right)^{\top}$ is close to $X X^{\top}$. Further, we'll give a sum-of-squares proof of this fact, which will give us an algorithm.

First, it will be useful for us to better understand the maximizer $Y^{*}$. The following characterization of the maximizer is easier to encode in a low-degree polynomial system.

Lemma 2.1. The balanced $k$-way partition $Y^{*}$ maximizing $\operatorname{Pr}[G \mid Y]$ is also the balanced $k$-way partition maximizing the function $\left\langle A_{G}, Y Y^{\top}\right\rangle \cdot \log \frac{p_{\text {in }}\left(1-p_{\text {out }}\right)}{p_{\text {out }}\left(1-p_{\text {in }}\right)}$, or equivalently,

$$
Y^{*}=\left\{\begin{array}{ll}
\arg \max _{Y}\left\langle A_{G}, Y Y^{\top}\right\rangle & \frac{p_{\text {in }}}{1-p_{\text {in }}}>\frac{p_{\text {out }}}{1-p_{\text {out }}} \\
\arg \min _{Y}\left\langle A_{G}, Y Y^{\top}\right\rangle & \frac{p_{\text {in }}}{1-p_{\text {in }}}<\frac{p_{\text {out }}}{1-p_{\text {out }}}
\end{array} .\right.
$$

Proof. For $i \in[n]$, let $Y_{i} \in\{0,1\}^{k}$ be such that $Y_{i}(c)=Y_{i c}$. Note that for $i, j \in[n],\left\langle Y_{i}, Y_{j}\right\rangle=1[i, j$ in same community $]$. Now since the edges of $G$ are independent conditioned on $Y$, we can write,

$$
\operatorname{Pr}[G \mid Y]=\prod_{i, j \in[n]}\left(p_{\text {in }}^{\left\langle Y_{i}, Y_{j}\right\rangle} \cdot p_{\text {out }}^{1-\left\langle Y_{i}, Y_{j}\right\rangle}\right)^{1[(i, j) \in E(G)]}\left(\left(1-p_{\text {in }}\right)^{\left\langle Y_{i}, Y_{j}\right\rangle}\left(1-p_{\text {out }}\right)^{1-\left\langle Y_{i}, Y_{j}\right\rangle}\right)^{1[(i, j) \notin E(G)]}
$$

Because $\log$ is a strictly increasing function, the maximizer of $\operatorname{Pr}[G \mid Y]$ is the same as the maximizer of $\log \operatorname{Pr}[G \mid Y]$. Now, applying the logarithm and simplifying the above, we can write,

$$
\begin{aligned}
\log \operatorname{Pr}[G \mid Y]= & \sum_{(i, j) \in E(G)}\left\langle Y_{i}, Y_{j}\right\rangle \log \frac{p_{\text {in }}}{p_{\text {out }}}+\sum_{(i, j) \notin E(G)}\left\langle Y_{i}, Y_{j}\right\rangle \log \frac{1-p_{\text {in }}}{1-p_{\text {out }}} \\
& +|E(G)| \log p_{\text {out }}+\left(\binom{n}{2}-|E(G)|\right) \log \left(1-p_{\text {out }}\right)
\end{aligned}
$$

The quantity on the second line does not depend on $Y$ at all, and so we can call it stuff $(G)$. Expressing the above as a matrix-matrix inner product,

$$
=\frac{1}{2} \log \frac{1-p_{\text {in }}}{1-p_{\text {out }}} \cdot\left\langle Y Y^{\top}, J\right\rangle+\frac{1}{2} \log \frac{p_{\text {in }}\left(1-p_{\text {out }}\right)}{p_{\text {out }}\left(1-p_{\text {in }}\right)} \cdot\left\langle Y Y^{\top}, A_{G}\right\rangle+\operatorname{stuff}(G)
$$

for $J$ the $n \times n$ all-1's matrix, and $A_{G}$ the adjacency matrix of $G .{ }^{1}$ We can simplify even further with the following claim:

[^0]Claim 2.2. For any balanced $k$-way partition $Y$ of $[n],\left\langle Y Y^{\top}, J\right\rangle=\frac{n^{2}}{k}$.
Proof of Claim 2.2. We can manipulate the expression algebraically, to start:

$$
\left\langle Y Y^{\top}, J\right\rangle=\left\langle Y Y^{\top}, 11^{\top}\right\rangle=\operatorname{Tr}\left(Y Y^{\top} 11^{\top}\right)=\left\|1^{\top} Y\right\|_{2}^{2} .
$$

Now, for each column $Y^{c}$ of $Y$, we have that $\left\langle 1, Y^{c}\right\rangle=\frac{n}{k}$ by the balanced assumption. This yields $\left\langle Y Y^{\top}, J\right\rangle=$ $\left(\frac{n}{k}\right)^{2} \cdot k$ as desired.

Applying Claim 2.2, we can push the $\left\langle Y Y^{\top}, J\right\rangle$ term into stuff $(G)$, creating a term stuff ${ }^{\prime}(G)$ which again does not depend on the specific choice of balanced partition $Y$. So,

$$
\log \operatorname{Pr}[G \mid Y]=\log \frac{p_{\text {in }}\left(1-p_{\text {out }}\right)}{p_{\text {out }}\left(1-p_{\text {in }}\right)} \cdot\left\langle Y Y^{\top}, A_{G}\right\rangle+\operatorname{stuff}^{\prime}(G),
$$

from which the conclusion follows.

### 2.1 Algorithm from the SoS paradigm

We will write a polynomial system which searches for $Y^{*}$. Our variables will be $Y \in \mathbb{R}^{n \times k}$, and our axioms will enforce that $Y$ corresponds to a balanced $k$-way partition:

$$
\mathcal{A}=\left\{Y_{i c}^{2}=Y_{i c}\right\}_{i \in[n], c \in[k]} \cup\left\{\left(1_{n}^{\top} Y\right)_{c}=\frac{n}{k}\right\}_{c \in[k]} \cup\left\{\left\langle Y_{i}, 1_{k}\right\rangle=1\right\}_{i \in[n]} .
$$

Our polynomial optimization program will be

$$
\max \left\langle Y Y^{\top}, A_{G}\right\rangle \text { s.t. } \mathcal{A} .
$$

The theorem below is not optimal in its dependence on $k$; see [HWX16, Ban18] for a sharper analysis of the performance of SDPs for exact recovery in block models with $k=2$, and [AS15] for an informationtheoretically optimal efficient algorithm.

Theorem 2.3. Suppose $p_{\text {in }}=\alpha \frac{\log n}{n}$, $p_{\text {out }}=\frac{\beta \log n}{n}$ satisfy $p_{\text {in }}>p_{\text {out }}{ }^{2}$, for $\alpha, \beta=\Omega(1)$. Then with high probability over the choice of $G$,

$$
\mathcal{A} \vdash_{2}\left\langle\frac{\widetilde{\mathrm{E}}\left[Y Y^{\top}\right]}{\left\|\widetilde{\mathrm{E}}\left[Y Y^{\top}\right]\right\|_{F}}, \frac{X X^{\top}}{\left\|X X^{\top}\right\|_{F}}\right\rangle \geqslant 1-\frac{\sqrt{k \alpha}+\sqrt{k^{3} \beta}}{\alpha-\beta} \cdot \sqrt{\frac{k}{\log n}} .
$$

This is to say that, if $\alpha$ and $\beta$ are sufficiently separated compared to $\sqrt{k \alpha}+\sqrt{\beta}$ and $\frac{k}{\log n}$, we know that $\widetilde{\mathrm{E}}\left[Y Y^{\top}\right]$ must be closely aligned with the block matrix $X X^{\top}$ which defines the communities, so the communities can be recovered up to some error (depending on $\varepsilon$ ). With more work one can actually show that an SDP-based algorithm recovers $X$ with no error with high probability; again see [HWX16, Ban18].

Proof. We want to use the fact that the objective value, $\left\langle Y Y^{\top}, A_{G}\right\rangle$ is large to conclude that $Y Y^{\top}$ should be close to $X X^{\top}$. To that end, we will split $A_{G}$ into a sum of its expectation and a random matrix representing its deviation,

$$
A_{G}=\mathrm{E}\left[A_{G}\right]+\Delta=p_{\text {out }} \cdot J_{n}+\left(p_{\text {in }}-p_{\text {out }}\right) \cdot X X^{\top}+\Delta .
$$

[^1]Now we can write,

$$
\begin{aligned}
\left\langle Y Y^{\top}, A_{G}\right\rangle & =p_{\text {out }} \cdot\left\langle Y Y^{\top}, J\right\rangle+\left(p_{\text {in }}-p_{\text {out }}\right)\left\langle Y Y^{\top}, X X^{\top}\right\rangle+\left\langle Y Y^{\top}, \Delta\right\rangle \\
& \leqslant p_{\text {out }} \cdot\left\langle Y Y^{\top}, J\right\rangle+\left(p_{\text {in }}-p_{\text {out }}\right)\left\langle Y Y^{\top}, X X^{\top}\right\rangle+\sum_{c \in[k]}\left\|Y^{c}\right\|^{2} \cdot \lambda_{\max }(\Delta) \\
& =p_{\text {out }} \cdot \frac{n^{2}}{k}+\left(p_{\text {in }}-p_{\text {out }}\right)\left\langle Y Y^{\top}, X X^{\top}\right\rangle+n \cdot \lambda_{\max }(\Delta),
\end{aligned}
$$

where the inequality was a degree- 2 sum-of-squares inequality (since $\Delta \leqslant \lambda_{\max }(\Delta) \cdot \mathbb{1}$ ) and the final equality uses $\mathcal{A}$ to conclude that $\left\|Y^{c}\right\|^{2}=\left\langle Y^{c}, 1_{n}\right\rangle=\frac{n}{k}$ for each $c \in[k]$, and also since the proof of Claim 2.2 is an SoS proof modulo the axioms $1^{\top} Y^{c}=\frac{n}{k}$ in $\mathcal{A}$. So we conclude that

$$
\begin{equation*}
\mathcal{A} \vdash_{2}\left(p_{\text {in }}-p_{\text {out }}\right)\left\langle Y Y^{\top}, X X^{\top}\right\rangle \geqslant\left\langle Y Y^{\top}, A_{G}\right\rangle-p_{\text {out }} \frac{n^{2}}{k}-n \lambda_{\max }(\Delta) . \tag{1}
\end{equation*}
$$

Now, we want to understand what this implies about $\widetilde{\mathrm{E}}\left[Y Y^{\top}\right]$ and $X X^{\top}$. Clearly $\mathcal{A}$ is feasible (so long as $\frac{n}{k}$ is an integer), so a pseudoexpectation operator respecting $\mathcal{A}$ must exist. Further, since the pseudoexpectation is a relaxation of a moment operator over the set of solutions to $\mathcal{A}$, we have that the maximizing (recall $p_{\text {in }}>p_{\text {out }}$ ) pseudoexpectation satisfies

$$
\begin{equation*}
\left\langle\widetilde{\mathbf{E}}\left[Y Y^{\top}\right], A_{G}\right\rangle \geqslant\left\langle X X^{\top}, A_{G}\right\rangle \approx \mathbf{E}_{G \mid X}\left[\left\langle X X^{\top}, A_{G}\right\rangle\right]=p_{\text {in }} \cdot \frac{n^{2}}{k} \tag{2}
\end{equation*}
$$

since $\left\langle X X^{\top}, A_{G}\right\rangle$ is a sum of independent random variables conditioned on $X$, so we expect it to concentrate quite well around its mean (so long as $p_{\mathrm{in}} n^{2} / k \gg 1$ ). We'll content ourselves with pretending this $\approx$ is an equality, keeping in mind that it is actually only true up to ( $1 \pm o(1)$ ) factors.

Hence combining (1) and (2),

$$
\begin{equation*}
\left\langle\widetilde{\mathrm{E}}\left[Y Y^{\top}\right], X X^{\top}\right\rangle \geqslant \frac{n^{2}}{k}-\frac{n \lambda_{\max }(\Delta)}{p_{\text {in }}-p_{\text {out }}} \tag{3}
\end{equation*}
$$

Also, since $X X^{\top}$ is a $0 / 1$ valued matrix, $\left\|X X^{\top}\right\|_{F}^{2}=\left\langle J, X X^{\top}\right\rangle=\frac{n^{2}}{k}$ (the final equality is by Claim 2.2). Our axioms imply, $\mathcal{A} \vdash_{2}\left\|Y Y^{\top}\right\|_{F}^{2}=\left\langle J, Y Y^{T}\right\rangle=\frac{n^{2}}{k}$, since the axioms $Y_{i c}^{2}=Y_{i c}$ and $\left\langle Y_{i}, 1_{k}\right\rangle=1$ imply that $\frac{n^{2}}{k}=$ $\left\langle J, Y Y^{\top}\right\rangle=\left\|Y Y^{\top}\right\|_{F}^{2}$. Since $\widetilde{\mathrm{E}}\left[a^{2}\right] \geqslant \widetilde{\mathrm{E}}[a]^{2}$ is a degree-2 sum-of-squares inequality, $\left\|\widetilde{\mathrm{E}}\left[Y Y^{\top}\right]\right\|_{F}^{2} \leqslant \widetilde{\mathrm{E}}\left\|Y Y^{\top}\right\|_{F}^{2}=\frac{n^{2}}{k}$. Using this information with (3),

$$
\left\langle\frac{\widetilde{\mathrm{E}}\left[Y Y^{\top}\right]}{\left\|\widetilde{\mathbb{E}}\left[Y Y^{\top}\right]\right\|_{F}}, \frac{X X^{\top}}{\left\|X X^{\top}\right\|_{F}}\right\rangle \geqslant 1-\frac{k \lambda_{\max }(\Delta)}{n\left(p_{\text {in }}-p_{\text {out }}\right)} .
$$

So we get some bound on the error, in terms of $\lambda_{\max }(\Delta)$.
What is $\lambda_{\text {max }}(\Delta)$ ? $\Delta$ is a block matrix with $k \times k$ blocks $\left\{\Delta_{a, b}\right\}_{a, b \in[k]}$ of size $\frac{n}{k} \times \frac{n}{k}$ each, where the blocks are indexed by pairs of communities. the entries of $\Delta_{a, b}$ are independent (up to symmetry, $\Delta_{a, b}=\Delta_{b, a}^{\top}$ ), each entry distributed like $\operatorname{Ber}\left(p_{\text {in }}\right)-p_{\text {in }}$ if $a=b$ and like $\operatorname{Ber}\left(p_{\text {out }}\right)-p_{\text {out }}$ if $a \neq b$. If the entries were all identically distributed as $\operatorname{Ber}(p)-p$ with $p=\Omega\left(\frac{\log n}{n}\right)$, we would have that $\|\Delta\| \leqslant 2 \sqrt{p(1-p) n}=$ $(1+o(1)) \cdot 2 \sqrt{\operatorname{Var}(\operatorname{Ber}(p)) n}$ with high probability. To get a very loose bound, we can write $\Delta$ as a sum of its blocks and apply the triangle inequality for the operator norm and get that with high probabiity,

$$
\lambda_{\max }(\Delta) \leqslant\|\Delta\| \leqslant \sum_{a, b \in[k]}\left\|\Delta_{a, b}\right\| \leqslant k \cdot \sqrt{p_{\text {in }}\left(1-p_{\text {in }}\right) \frac{n}{k}}+k(k-1) \cdot \sqrt{p_{\text {out }}\left(1-p_{\text {out }}\right) \frac{n}{k}} \leqslant \sqrt{p_{\text {in }} n k}+\sqrt{p_{\text {out }} n k^{3}} .
$$

This bound is not tight, but it is sufficient to get us something nontrivial. Hence, parametrizing $p_{\text {in }}=\alpha \frac{\log n}{n}$ and $p_{\text {out }}=\beta \frac{\log n}{n}$, we have

$$
\frac{k \lambda_{\max }(\Delta)}{n\left(p_{\text {in }}-p_{\text {out }}\right)} \leqslant \frac{k}{n\left(p_{\text {in }}-p_{\text {out }}\right)} \cdot\left(\sqrt{p_{\text {in }} n k}+\sqrt{p_{\text {out }} n k^{3}}\right)=\frac{\sqrt{k \alpha}+\sqrt{\beta k^{3}}}{(\alpha-\beta)} \cdot \sqrt{\frac{k}{\log n}} .
$$

This completes the proof.

## 3 An information-computation gap in the sparse regime

The problem of detection in stochastic block models has an information-computation gap when $p_{\text {in }}, p_{\text {out }}=$ $\Theta\left(\frac{1}{n}\right)$. Let $p_{\text {in }}=\frac{a}{n}$ and $p_{\text {out }}=\frac{b}{n}$, and denote also the average degree $d_{\text {avg }}=\frac{1}{k} a+\frac{k-1}{k} b$. As is intuitively clear, the problem of hypothesis testing between $\mathcal{H}_{0}=\mathcal{C}\left(n, p_{\text {avg }}\right)\left(\right.$ for $\left.p_{\text {avg }}=\frac{1}{k} p_{\text {in }}+\frac{k-1}{k} p_{\text {out }}\right)$ and $\mathcal{H}_{1}=$ $\mathcal{C}\left(n, k, p_{\text {in }}, p_{\text {out }}\right)$ gets easier as $p_{\text {in }}, p_{\text {out }}($ or $a, b)$ get further apart.

We'll associate the following signal-to-noise ratio parameter with this problem:

$$
s=\left(\frac{a-b}{k}\right)^{2} \cdot \frac{1}{d_{\mathrm{avg}}} .
$$

This may look unnatural at first, but below we'll explain where this quantity comes from.
The parameter setting where $s=1$ is called the Kesten-Stigum threshold (or KS-threshold). For all $k \geqslant 2$, the best detection algorithms known to date succeed if and only if $s>1$, or "above the KestenStigum threshold" [AS16a, BMR21].

In the special case $k=2$, the KS threshold $\tau_{\mathrm{KS}}=1$ and the information-theoretic threshold $\tau_{\text {info }}$ below which (when $\left.s<\tau_{\text {info }}\right) \mathrm{d}_{\mathrm{TV}}\left(\mathcal{H}_{0}, \mathcal{H}_{1}\right)=o(1)$ are known to coincide [MNS15, MNS18, Mas14]. However, the information theoretic threshold is known to scale with $k$ as $\tau_{\text {info }}=\Theta\left(\frac{\log k}{k}\right)$ [BMNN16]. Hence, for $k$ large enough, for $\Theta\left(\frac{\log k}{k}\right)<s \leqslant 1$, the detection problem is solvable information-theoretically, but we don't know computationally efficient algorithms that do so.

### 3.1 Heuristic justification for the KS threshold

The signal-to-noise parameter $s$ and the threshold $s=1$ come from an information-theoretic lower bound for solving the related root recovery problem in a broadcast process on a random tree, generated in such a manner that this random tree can be coupled with a breadth-first-search starting at an arbitrary vertex in $G \sim \mathcal{G}\left(n, k, p_{\text {in }}, p_{\text {out }}\right)$.

Breadth-first-search and/or a broadcast process on a tree. Imagine doing breadth-first search (BFS) up to depth $t$ starting at some node $v \in V(G)$. We can couple this BFS with the sampling of $G$ itself, for any constant $t$, as follows:

1. Start at the root, $v$.
2. Assign $v$ a community label $\in[k]$ uniformly at random.
3. Push $v$ onto a queue.
4. While the queue is non-empty, or until the vertex at the head of the queue is of distance $t$ from the root $v$ :
(a) Remove the leading vertex $u$ from the queue.
(b) Independently sample Pois $\left(\frac{1}{k} a\right)$ children of $u$ with the same label $c$ as $u$, and Pois $\left(\frac{1}{k} b\right)$ children with label $c^{\prime}$ for each $c^{\prime} \in[k] \backslash\{c\}$.
(c) Put each of these children into the queue.

One can show that for $t$ constant, this process can be coupled with a breadth-first search in $G$ which succeeds with high probability, by using the fact that $\operatorname{Bin}\left(n, \frac{\lambda}{n}\right)$ is well-approximated by $\operatorname{Pois}(\lambda)$ as $n \rightarrow \infty$. This is a classic style of coupling, see for example [AS16b], Chapter 11.

Now, this can also be viewed as a broadcast process on a Galton-Watson tree of depth at most $t$. The idea is that each node samples Pois $\left(d_{\mathrm{avg}}\right)$ children idependently, then "broadcasts" a label to each of its children, keeping its own label with probability $\frac{a}{d_{\text {avg }}}$ and choosing one of the $k-1$ other labels uniformly at random with the remaining probability. By Poisson splitting, this is equivalent to the process described above.

Root reconstruction and a heuristic lower bound. A natural question in the context of a broadcast process on a tree is the following: given the labels at the leaves of the tree, an we get a reasonable estimate for the label of the root? The KS threshold arises as the threshold at which root reconstruction becomes information-theoretically possible [KS66, EKPS00]. Below we'll prove one direction: we'll show that when $s>1$, root reconstruction is possible with a canonical estimator.

Theorem 3.1. Let $X_{t}$ be the number of vertices with labelc at deptht in the tree (so $X_{0}=1$ [root has label c]). Then

$$
\lim _{t \rightarrow \infty} \frac{\mathrm{E}\left[X_{t} \mid X_{0}=1\right]-\mathrm{E}\left[X_{t} \mid X_{0}=0\right]}{\sqrt{\operatorname{Var}\left[X_{t}\right]}}=\frac{1}{k} s^{t / 2}
$$

In particular if $s>1$, one can use the value of $X_{t}$ for $t$ large enough to estimate whether $X_{0}=1$ (equivalently, if the root has label $c$ ) by checking if $X_{t}$ is closer to $\mathrm{E}\left[X_{t} \mid X_{0}=1\right]$ or $\mathrm{E}\left[X_{t} \mid X_{0}=0\right]$, with the bound on the variance furnishing concentration via the second moment method. When $s \leqslant 1$, the difference in expectations is overwhelmed by the variance.

Before we dive into the proof, what does root reconstruction have to do with detection in block models? This has to do with the performance of the Belief Propagation Algorithm (BP), which precisely computes the marginal distribution of the root vertex in a broadcast process on a tree (given access to the leaf vertex labels, and to the structure of the tree). If we are in the regime $s>1$ where root reconstruction is possible, then Belief Propagaion also works for reconstructing the root, and in fact it achieves the statistically optimal estimation rate.

Though Belief Propagation is designed to work on trees, people can (and do) run it on graphs with cycles; it seems to work okay on sparse graphs, and in some cases one can even prove formally that some version of it works [KMM ${ }^{+} 13$, AS16a, BLM15]. The reason $s>1$ might suggest that BP works is that if you choose a random partition of the vertices of $G$ to start, then patch it up using BP as though each vertex were the root of a depth- $t$ tree (in sparse random graphs, most vertices' depth- $t$ neighborhoods are trees), you get nontrivial information about some fraction of the vertices' labels. Alternatively if $s \leqslant 1$ and BP fails, then this could be taken as evidence that any polynomial-time algorithm will fail, if you beleive that BP is optimal. This is discussed further in, for example, these sources [DKMZ11, M ${ }^{+}$17, LM21].

It is surprising that such imprecise heuristic reasoning seems to predict the limits of performance not only of BP algorithms, but also of dramatically different algorithms like SoS/SDPs [MS16, BMR21] and lowdegree polynomial estimators [HS17, $\mathrm{BBK}^{+} 21$ ]. This apparent universality across algorithms is a theme we'll return to throughout the class.

Now, we prove that $X_{t}$ is a reasonable estimator for whether $X_{0}=1$.

Proof of Theorem 3.1. Let $X_{t}$ be the number of vertices of label $c$ at depth $t$, and let $Y_{t}$ be the number of vertices of label not $c$ at depth $t$. By definition of our process,

$$
\begin{aligned}
\mathrm{E}\left[X_{t} \mid X_{t-1}, Y_{t-1}\right] & =\frac{1}{k} a \cdot X_{t-1}+\frac{1}{k} b \cdot Y_{t-1} \\
\mathrm{E}\left[Y_{t} \mid X_{t-1}, Y_{t-1}\right] & =\frac{k-1}{k} b \cdot X_{t-1}+\left(\frac{a}{b}+\frac{k-2}{k} b\right) \cdot Y_{t-1} .
\end{aligned}
$$

Encoding this linear-algebraically for convenience, we have that

$$
\mathbf{E}\left[\begin{array}{ll}
X_{t} & Y_{t}
\end{array}\right]=\left[\begin{array}{ll}
X_{t-1} & Y_{t-1}
\end{array}\right] A, \quad \text { for } \quad A=\left[\begin{array}{cc}
\frac{1}{k} a & \frac{k-1}{k} b \\
\frac{1}{k} b & \frac{1}{k} a+\frac{k-2}{k} b
\end{array}\right] .
$$

So $A$ is like the "transition matrix" for the expected number of vertices of each type, with the first row/column corresponding to $X$ and the second row/column corresponding to $Y$. One can check that $A$ has eigenvalues $\frac{a-b}{k}$ and $d_{\mathrm{avg}}=\frac{a+(k-1) b}{k}$, with corresponding left-eigenvectors $[1-1]$ and $[1 k-1]$ respectively.

Now, notice that

$$
E\left[X_{t} \mid X_{0}=1\right]-\mathbf{E}\left[X_{t} \mid X_{0}=0\right]=e_{1}^{\top} A^{t} e_{1}-e_{2}^{\top} A^{t} e_{1}=\left(e_{1}-e_{2}\right)^{\top} A^{t} e_{1}=\left(\frac{a-b}{k}\right)^{t}
$$

because $e_{1}-e_{2}$ is the left eigenvector of $A$ with eigenvalue $(a-b) / k$.
We now compute $\operatorname{Var}\left[X_{t}\right]$. This is made simple if we consider $\operatorname{Var}\left[X_{t} \mid X_{t-1}, Y_{t-1}\right]$. Conditioned on $X_{t-1}, Y_{t-1}, X_{t}$ is a sum of independent Poisson random variables, one for the number of children of community $c$ for each node in the $t-1$ th leve. By independence, the variance of the sum is simply the sum of the variances. Further, for $Q \sim \operatorname{Pois}(\lambda), \operatorname{Var}[Q]=\lambda$. This implies that

$$
\operatorname{Var}\left[X_{t} \mid X_{t-1}, Y_{t-1}\right]=\frac{1}{k} a X_{t-1}+\frac{1}{k} b Y_{t-1}=\mathbf{E}\left[X_{t} \mid X_{t-1}, Y_{t-1}\right],
$$

And hence, we can compute

$$
\operatorname{Var}\left[X_{t}\right]=\mathrm{E}\left[X_{t}\right]=\left[\begin{array}{cc}
\frac{1}{k} & \frac{k-1}{k}
\end{array}\right] A^{t} e_{1}=\frac{1}{k} d_{\mathrm{avg}}^{t}
$$

where we have used that $\operatorname{Pr}\left[X_{0}=1\right]=\frac{1}{k}$, and that $[1 k-1]$ is the eigenvector of $A$ corresponding to the eigenvalue $d_{\text {avg }}$.

Combining the above,

$$
\lim _{t \rightarrow \infty} \frac{\mathrm{E}\left[X_{t} \mid X_{0}=1\right]-\mathrm{E}\left[X_{t} \mid X_{0}=0\right]}{\sqrt{\operatorname{Var}\left[X_{t}\right]}}=\lim _{t \rightarrow \infty} \frac{1}{k}\left(\frac{a-b}{k \sqrt{d_{\mathrm{avg}}}}\right)^{t}
$$

and the quantity on the right can be re-written as $\frac{1}{k} s^{t / 2}$, as desired.

### 3.2 Weak recovery

In addition to the question of detection in the sparse regime, one can also ask about recovery. It's not hard to convince yourself that exact recovery is impossible, even when detection is possible: since $p_{\text {in }}, p_{\text {out }}=\Theta\left(\frac{1}{n}\right)$, a constant fraction of the vertices will be isolated, and there is no hope in recovering their community membership. Still, you could hope to recover the partitions with accuracy better than a uniform random assignment. This algorithmic task is called weak recovery. A very recent paper [YP22] gives the
information-theoretically optimal error rate for weak recovery everywhere above the KS-threshold, for the special case $k=2$.

An interesting open problem is whether this algorithm can be matched by SoS algorithms; one advantage of SoS algorithms and SDPs is that they typically enjoy strong robustness guarantees, see e.g. [DdNS22]. Robust SDP algorithms are known to achieve nontrivial weak recovery above the KS threshold, but it is open whether they achieve the optimal rates. The work of Banks, Mohanty, and Raghavendra [BMR21] presents an intriguing approach.

## 4 Conclusion

Bibliographic remarks. There is a vast body of literature on the stochastic block model, and I won't do it justice here; the surveys [Abb17, $\mathrm{M}^{+} 17$ ] are nice resources. See also the specific reference pointers throughout the notes above, though there are many references that I have left out.

Contact. Comments are welcome at tselil@stanford.edu.

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[^0]:    ${ }^{1}$ Really to be absolutely correct we should subtract the diagonal of $J$, since we don't consider the $i=j$ terms. However the diagonal of $Y Y^{\top}$ is just $\mathbb{1}$, so this will not matter, and you can check that this does not affect the overall argument.

[^1]:    ${ }^{2}$ The opposite case can be handled by a near-identical proof

