

(Nearly) Efficient Algorithms for the Graph Matching Problem

Tselil Schramm
(Harvard/MIT)

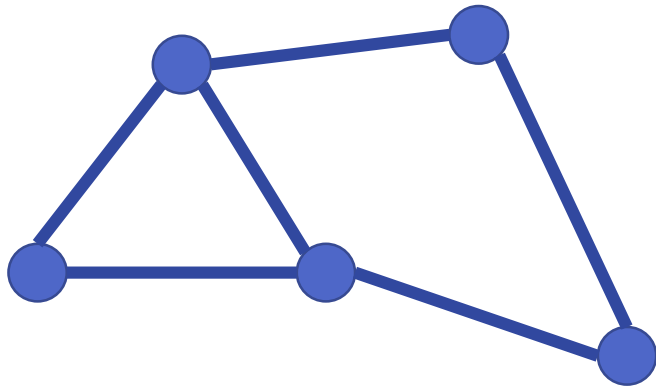
with Boaz Barak, Chi-Ning Chou, Zhixian Lei & Yueqi Sheng (Harvard)

graph matching problem (approximate graph isomorphism)

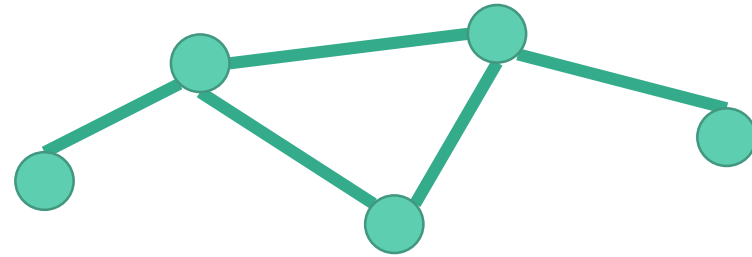
input: two graphs on n vertices

goal: find permutation of vertices that maximizes # shared edges

$$\max_{\pi} \langle A_{G_0}, \pi(A_{G_1}) \rangle$$



G_0



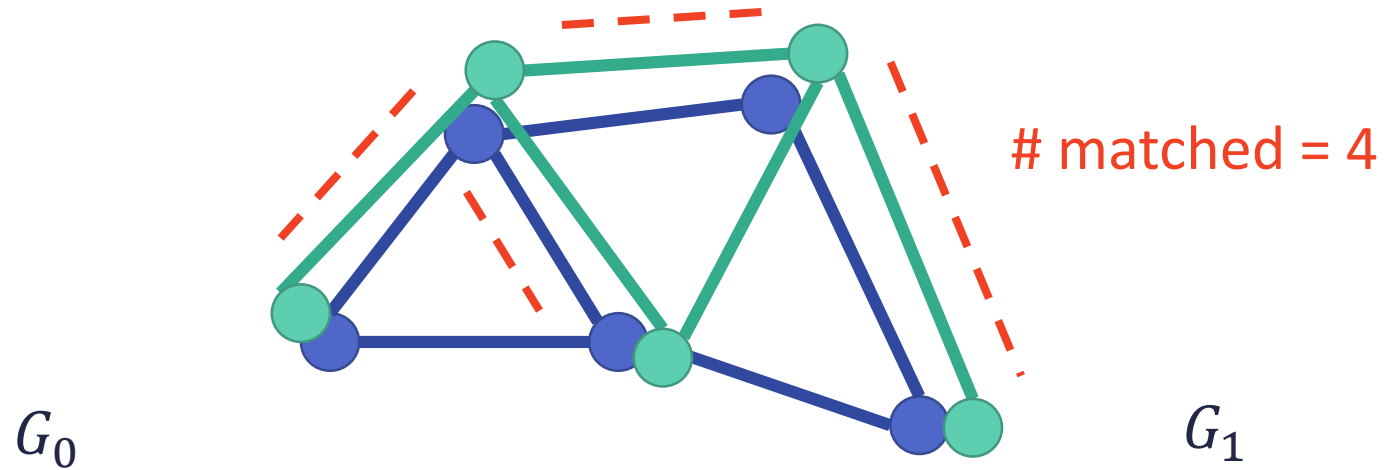
G_1

graph matching problem (approximate graph isomorphism)

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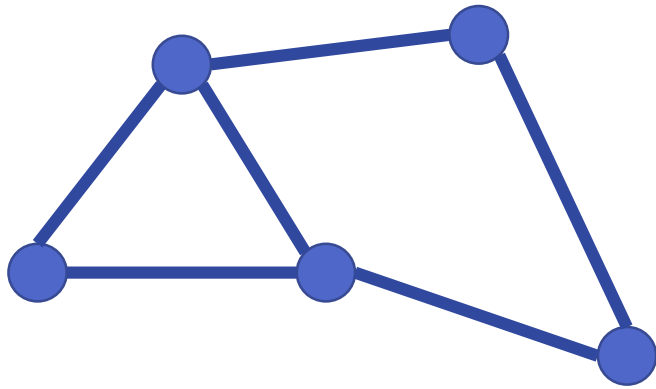


graph matching problem (approximate graph isomorphism)

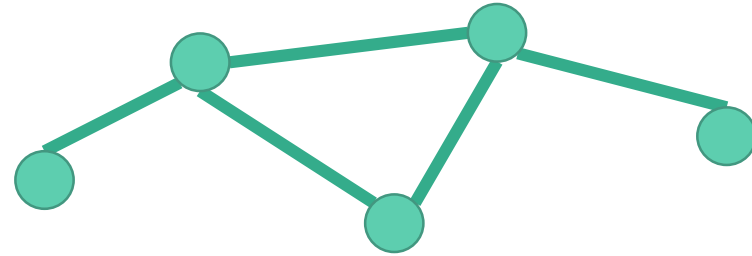
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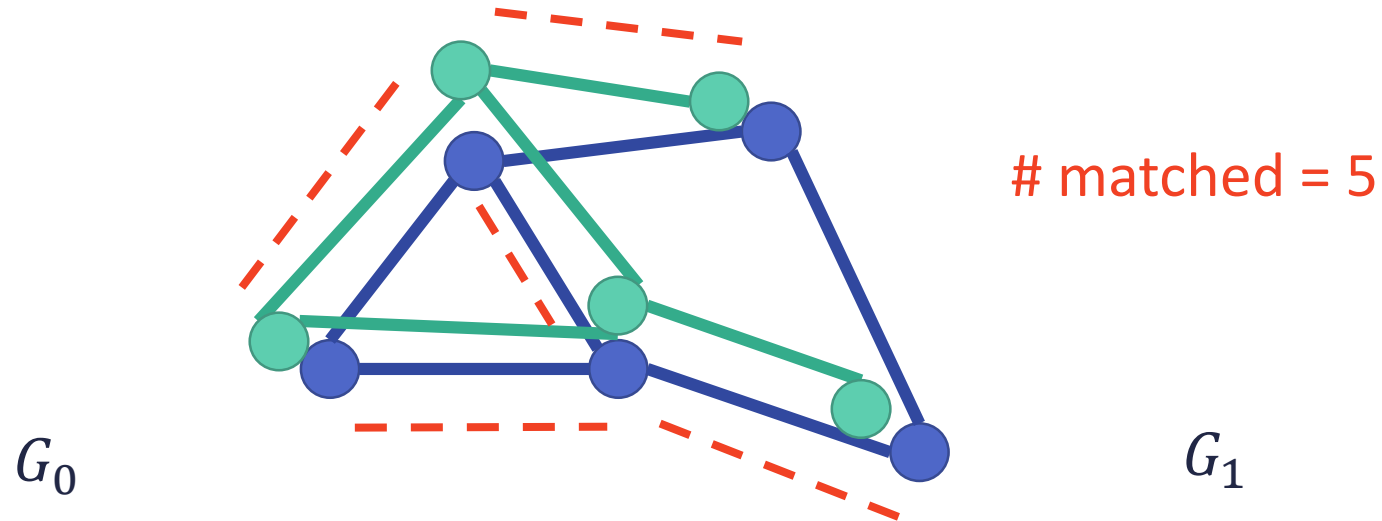
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graph matching problem (approximate graph isomorphism)

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computationally hard (of course)

NP-hard: reduction from quadratic assignment problem (non-simple graphs).

[Lawler'63]

also: reduction from sparse random 3-SAT to approximate version

[O'Donnell-Wright-Wu-Zhou'14]

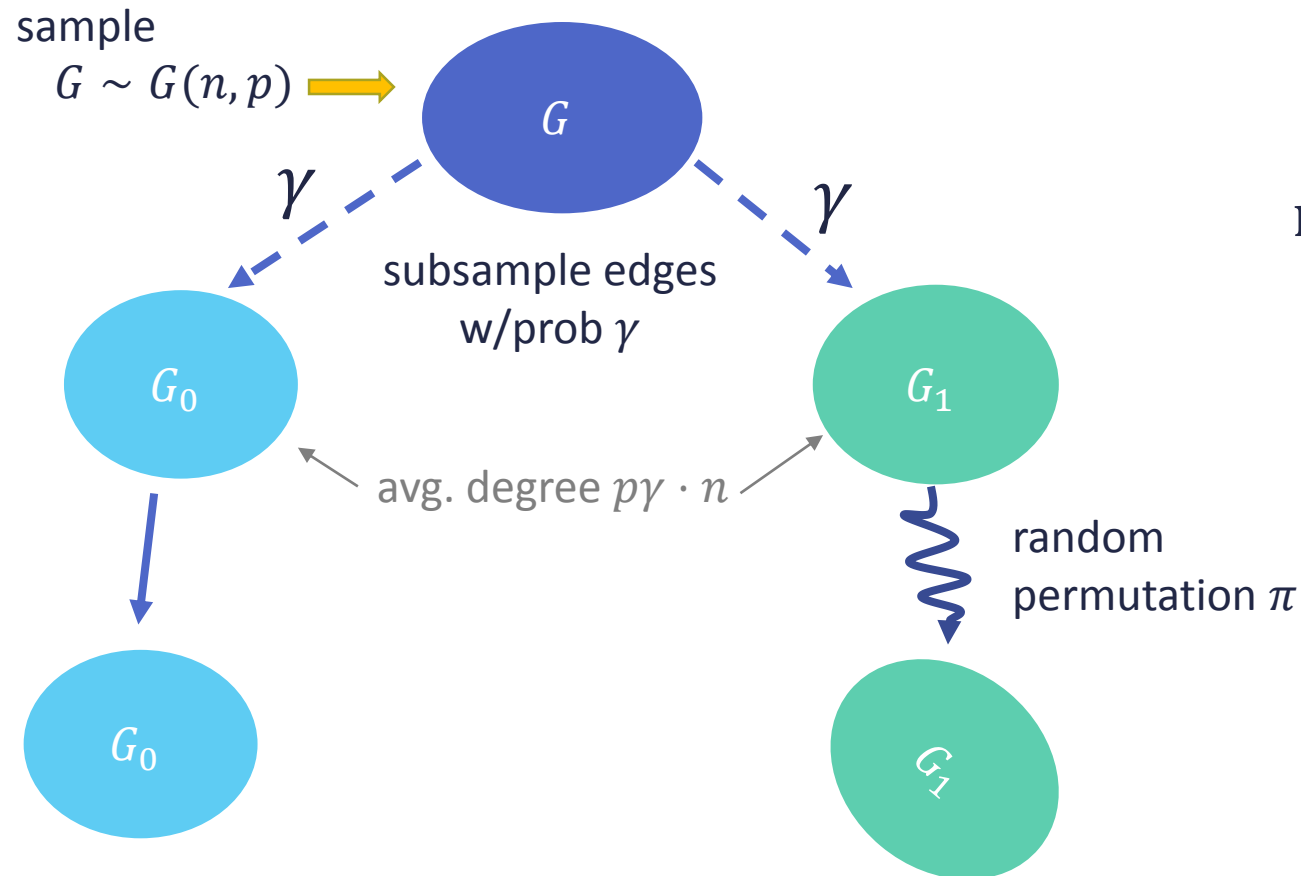
practitioners: undeterred

- computational biology [e.g. Singh-Xu-Berger'08]
- de-anonymization [e.g. Narayanan-Shmatikov'09]
- social networks [e.g. Korula-Lattanzi'14]
- image alignment [e.g. Cho-Lee'12]
- machine learning [e.g. Cour-Srinivasan-Shi'07]
- pattern recognition, e.g.
 - “thirty years of graph matching in pattern recognition”
[Conte-Foggia-Sansone-Vento'04]

“robust average-case graph isomorphism”

average case: correlated random graphs

structured model



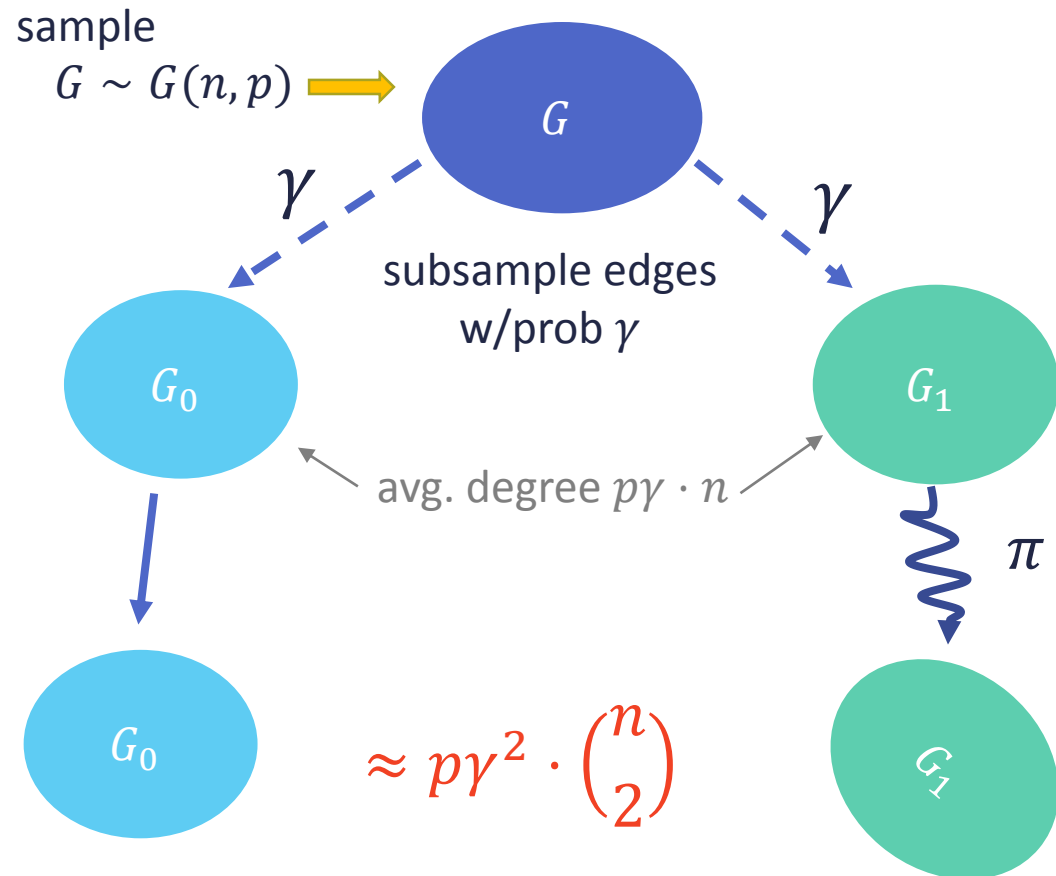
$$\max_{\pi} \langle A_{G_0}, \pi(A_{G_1}) \rangle \approx p\gamma^2 \cdot \binom{n}{2}$$

[e.g. Pedarsani-Grossglauser'11,
Lyzinski-Fishkind-Priebe'14,
Korula-Lattanzi'14]

“robust average-case graph isomorphism”

average case: correlated random graphs

structured model

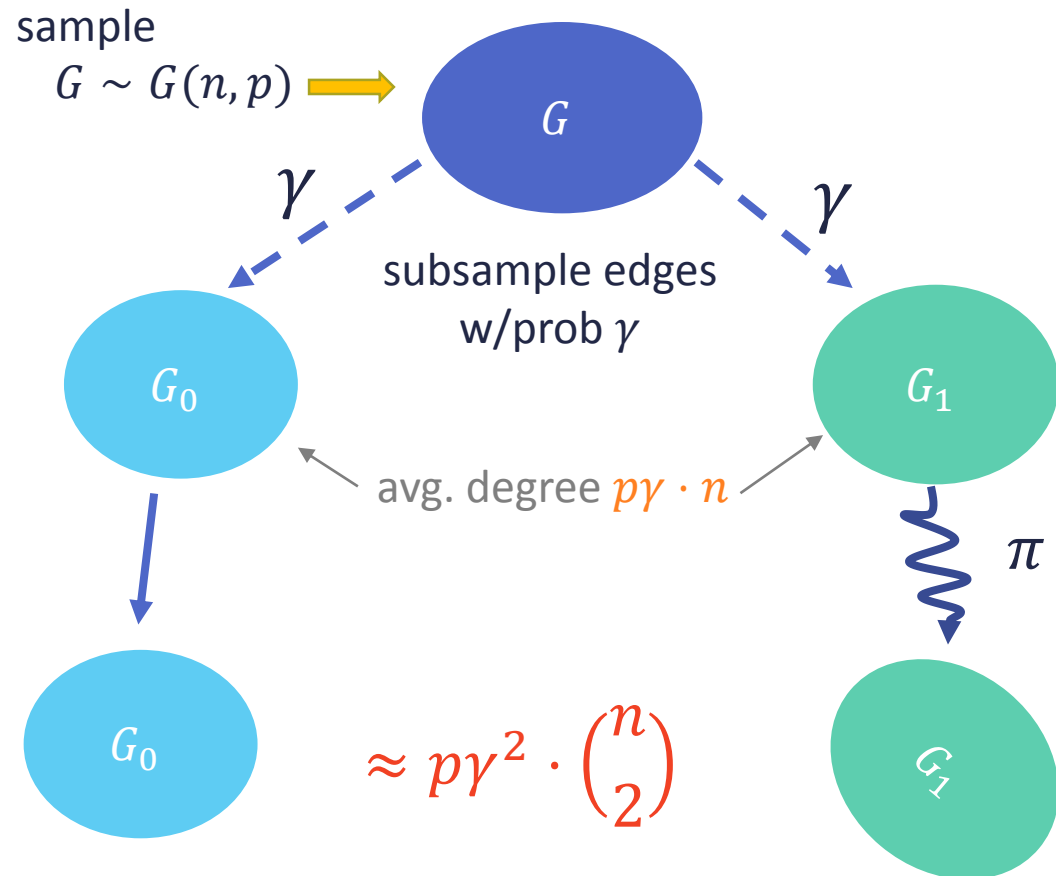


“null” model

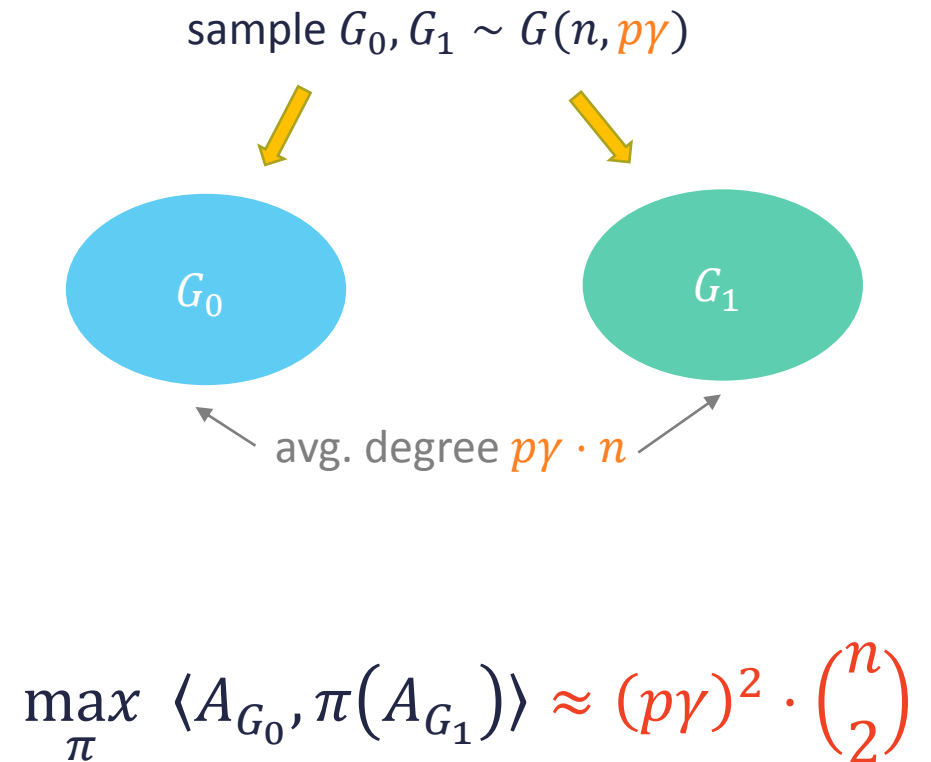
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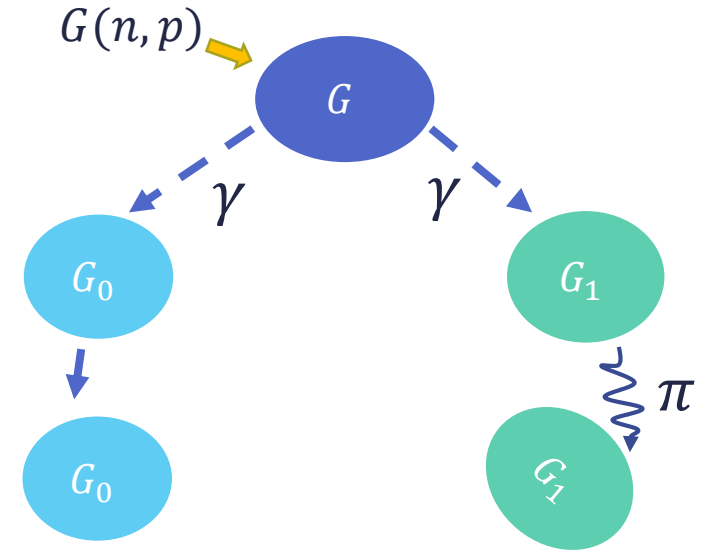


“null” model



information theoretic limit

for which p, γ can we recover π ?



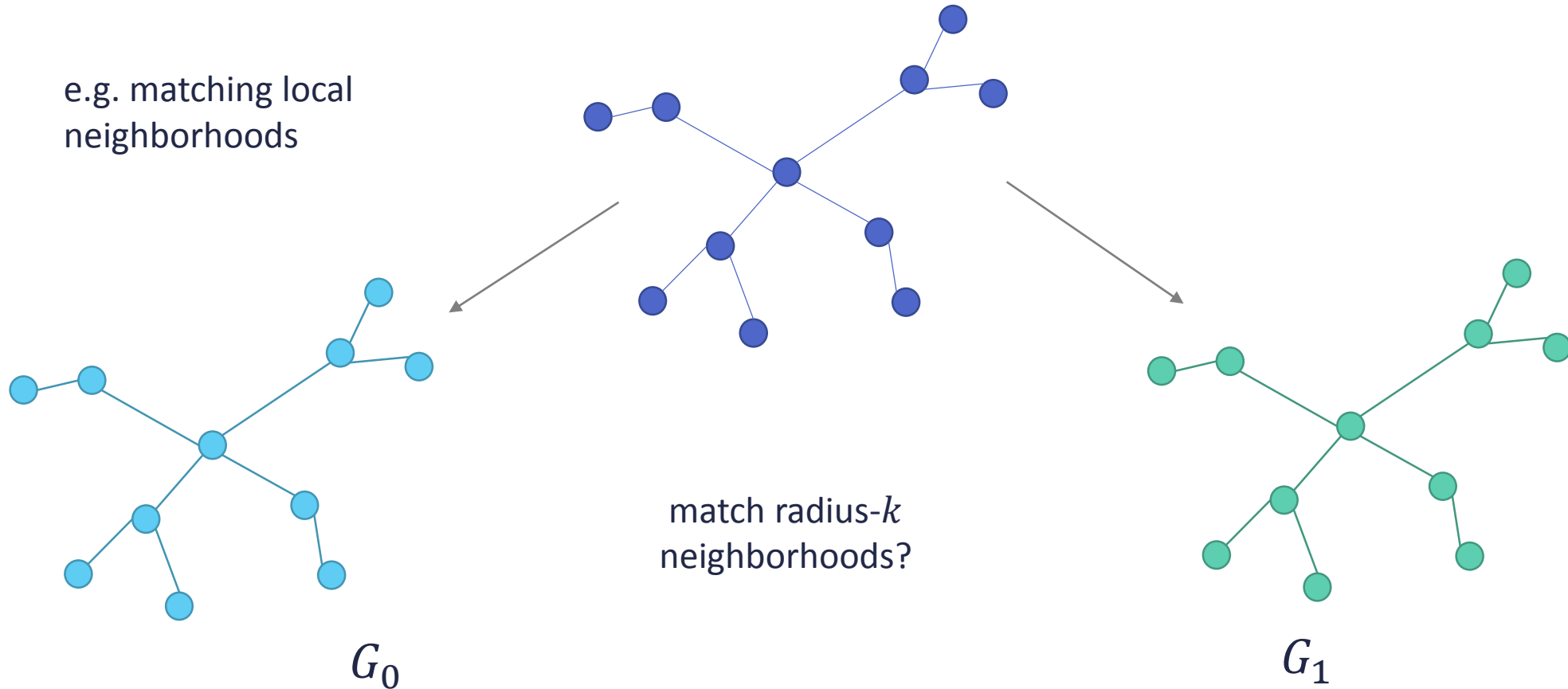
Theorem [Cullina-Kivayash'16&17]

Iff $p\gamma^2 > \frac{\log n}{n}$, with high probability π is the unique maximizing permutation.

algorithms for robust average case?

average-case graph isomorphism algorithms fail.

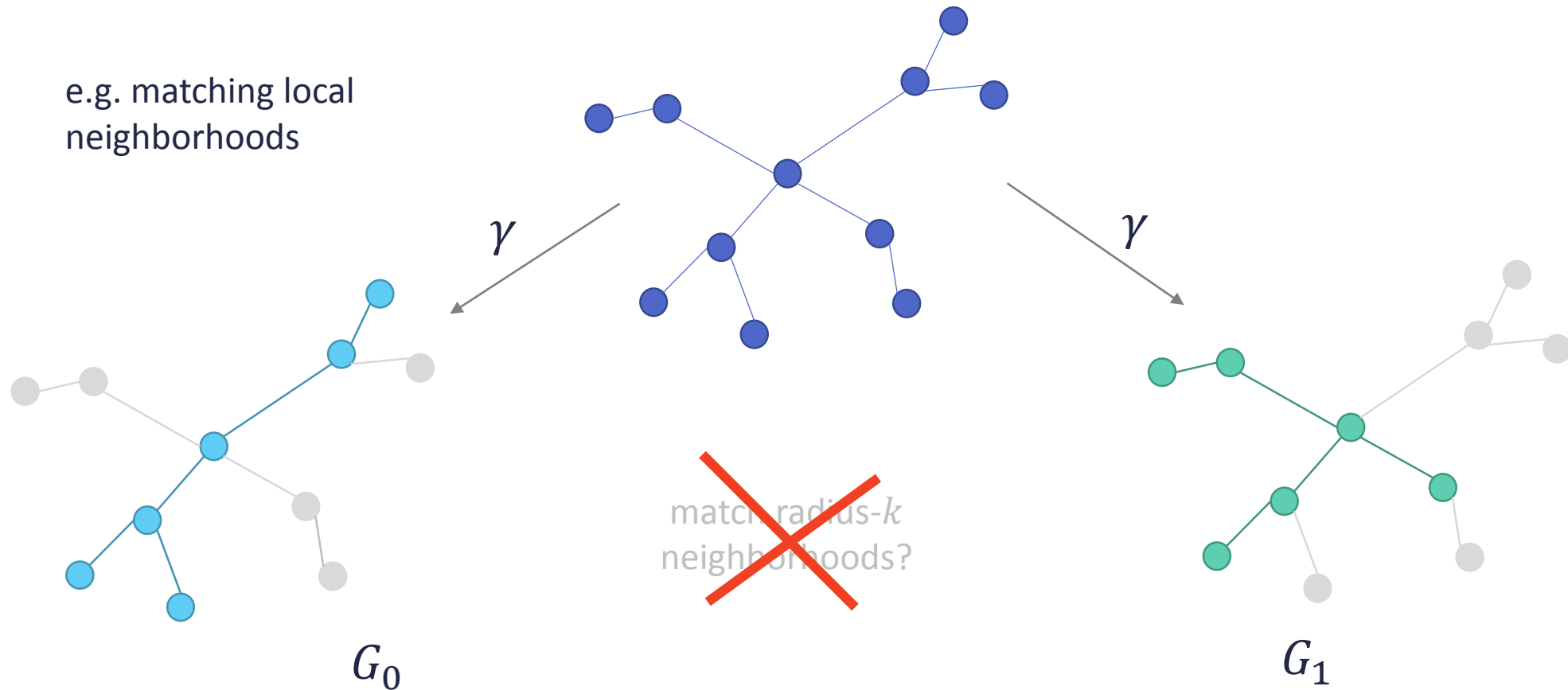
e.g. matching local neighborhoods



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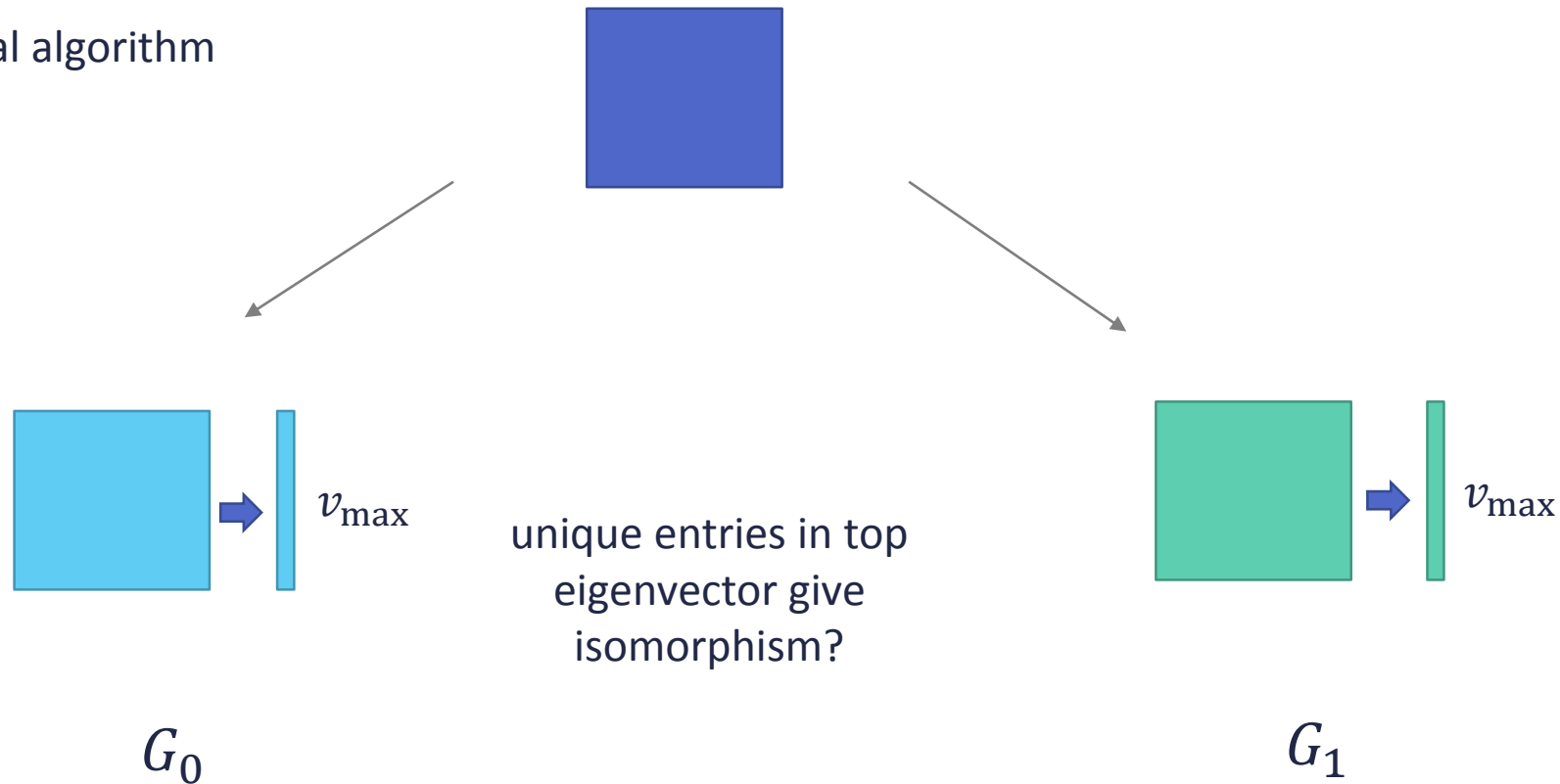
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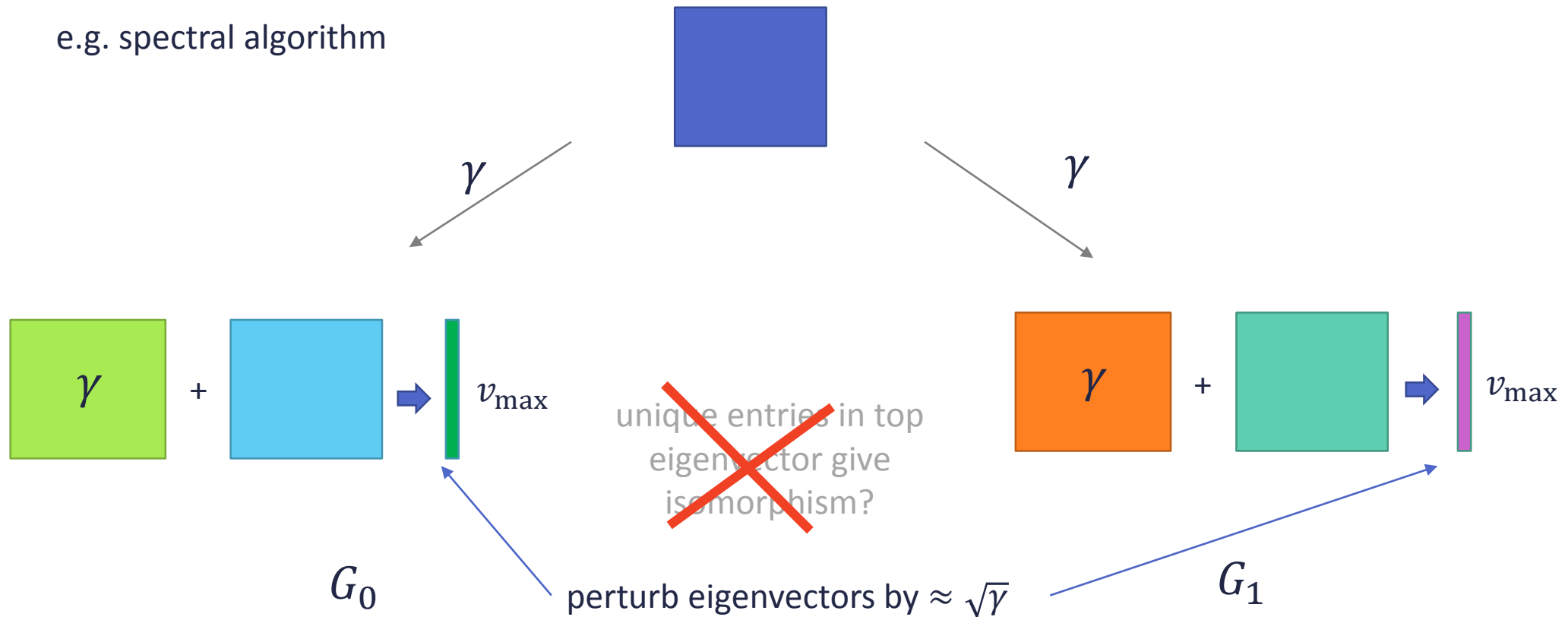
e.g. spectral algorithm



algorithms for robust average case?

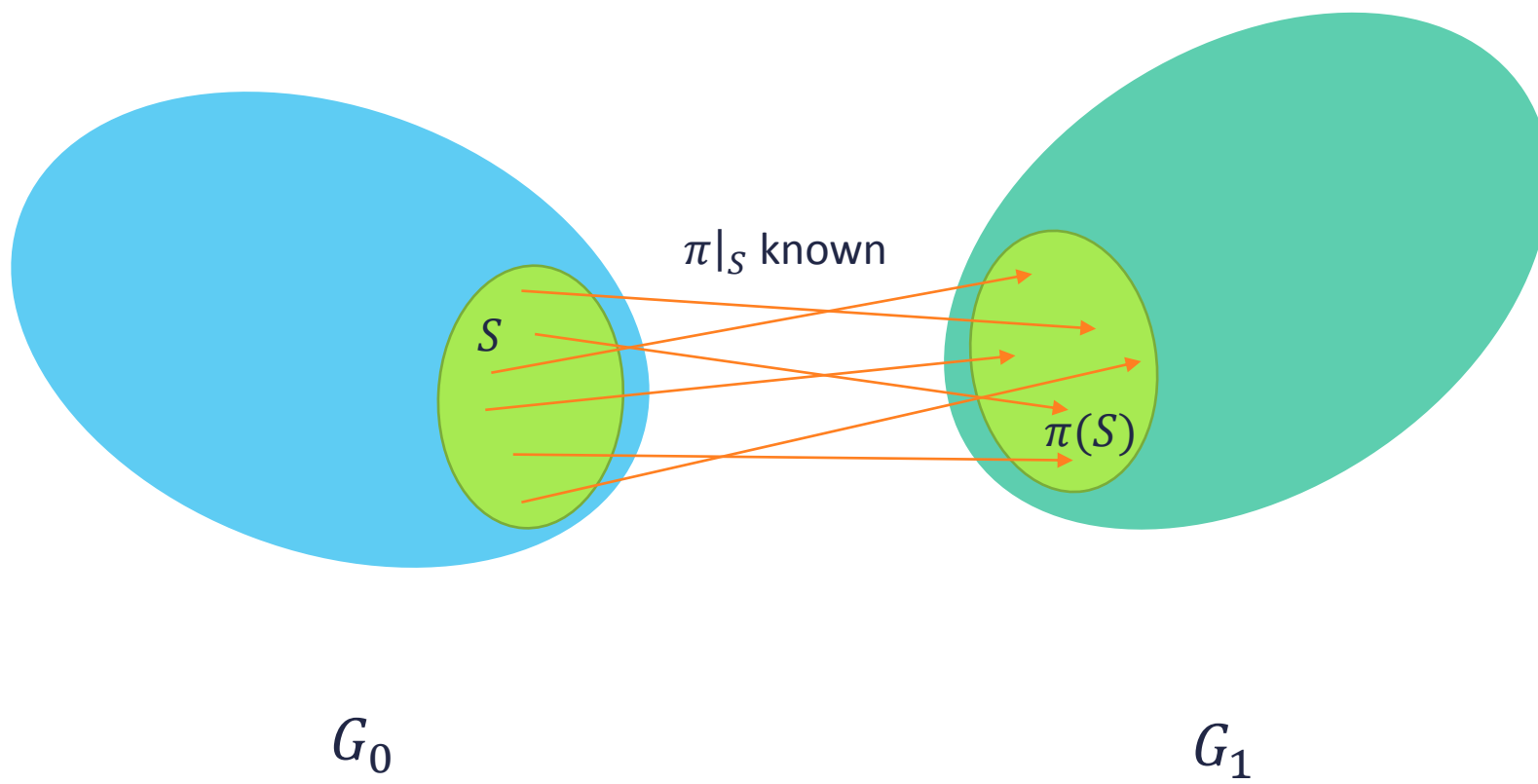
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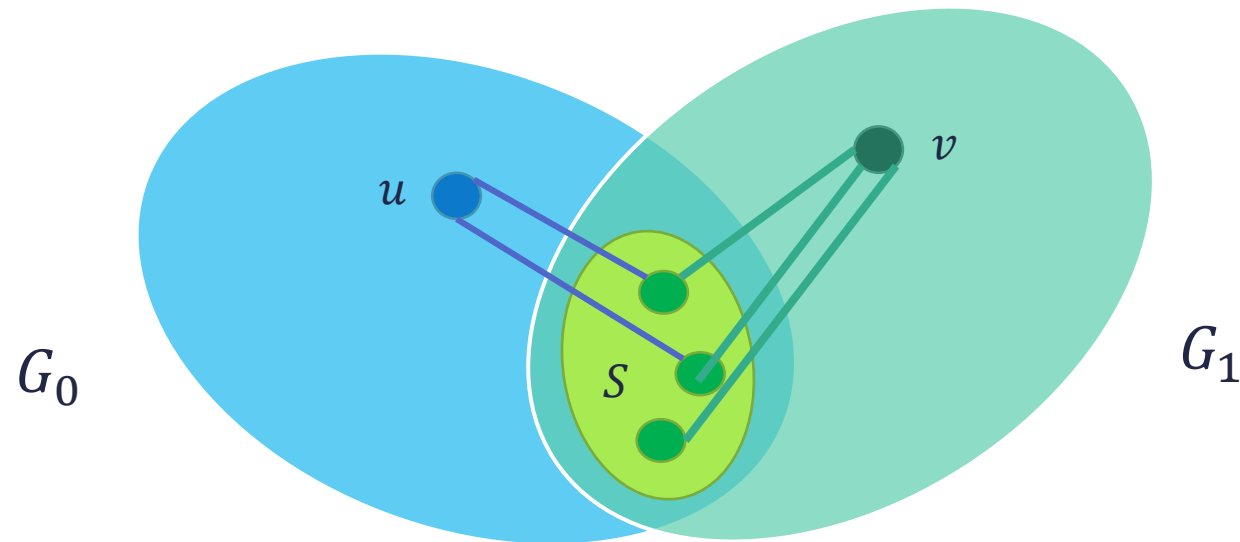
actual algorithms for robust average case?

starting from a seed



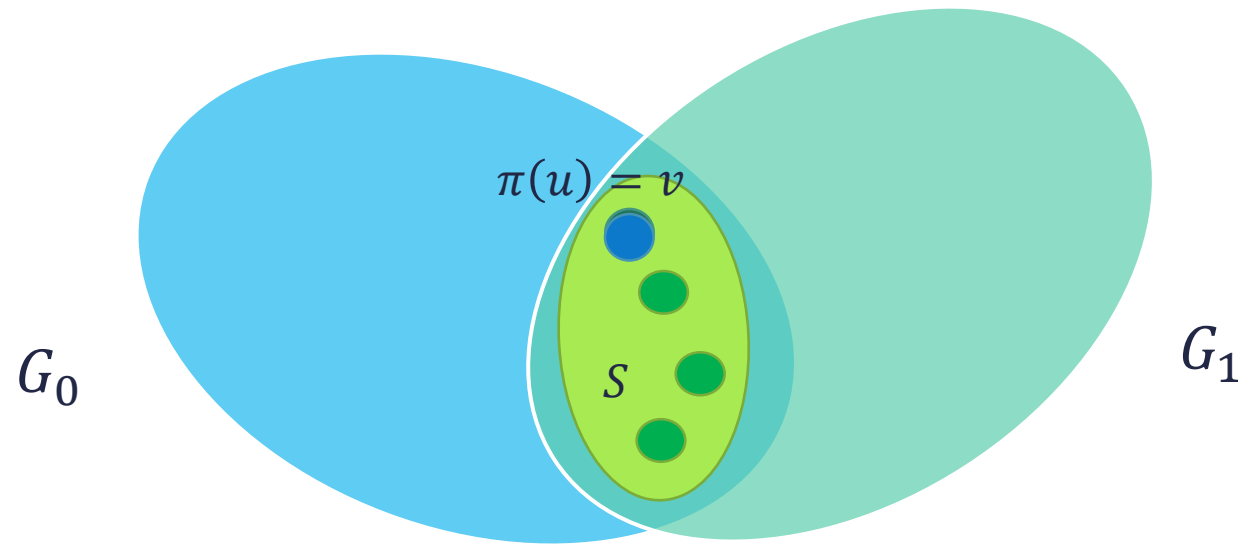
starting from a seed

match vertices with similar adjacency into S



starting from a seed

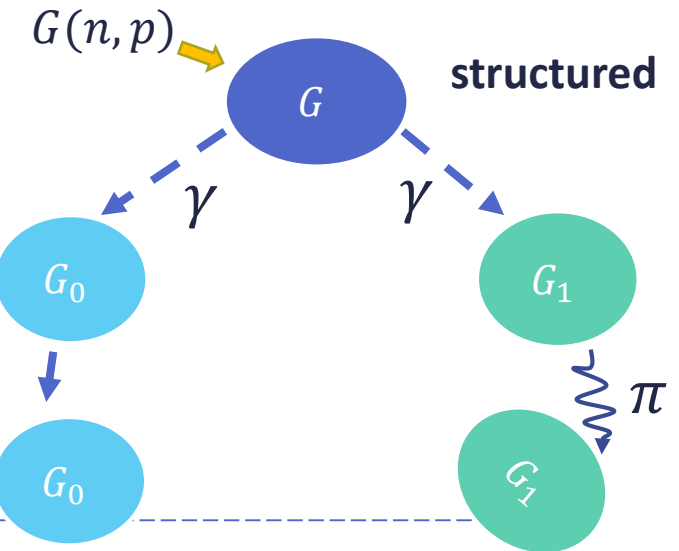
match vertices with similar adjacency into S



iff $\text{seed} \geq \Omega(n^\epsilon)$, the seeded algorithm approximately recovers π . [Yartseva-Grossglauser'13]

need $2^{\tilde{O}(n^\epsilon)}$ time to guess a seed.

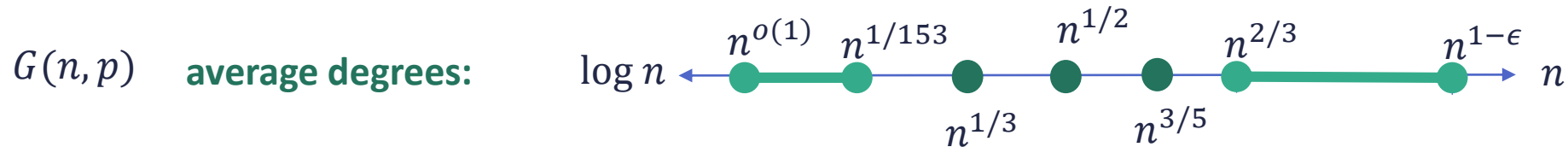
our results



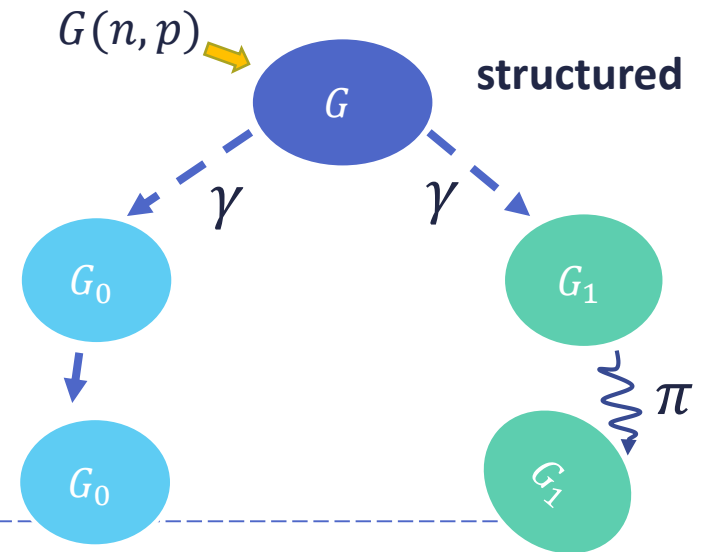
Theorem

For any $\epsilon > 0$, if $p \in \left[\frac{n^{o(1)}}{n}, \frac{n^{\frac{1}{153}}}{n} \right] \cup \left[\frac{n^{\frac{2}{3}}}{n}, \frac{n^{1-\epsilon}}{n} \right]$ and $\gamma = \Omega(1)$,* there is a $n^{O(\log n)}$ time algorithm that recovers π on $n - o(n)$ of the vertices w/prob ≥ 0.99 .

*we allow $\gamma = \Omega\left(\frac{1}{\log \log n}\right)$



our results



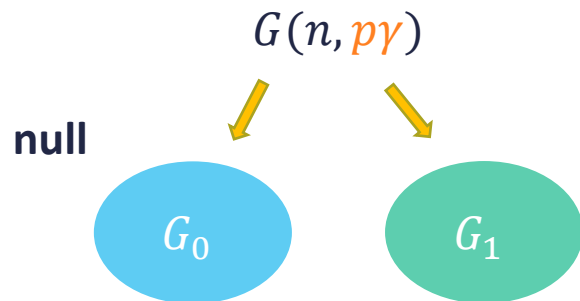
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*we allow $\gamma \geq \frac{1}{\log^{o(1)} n}$

Theorem

If p, γ are as above then there is a $\text{poly}(n)$ time *distinguishing* algorithm for the **structured** vs **null** distributions.



hypothesis testing

our approach: small subgraphs

hypothesis testing: correlation of subgraph counts

recovery: match rare subgraphs

seedless algorithms!

outline

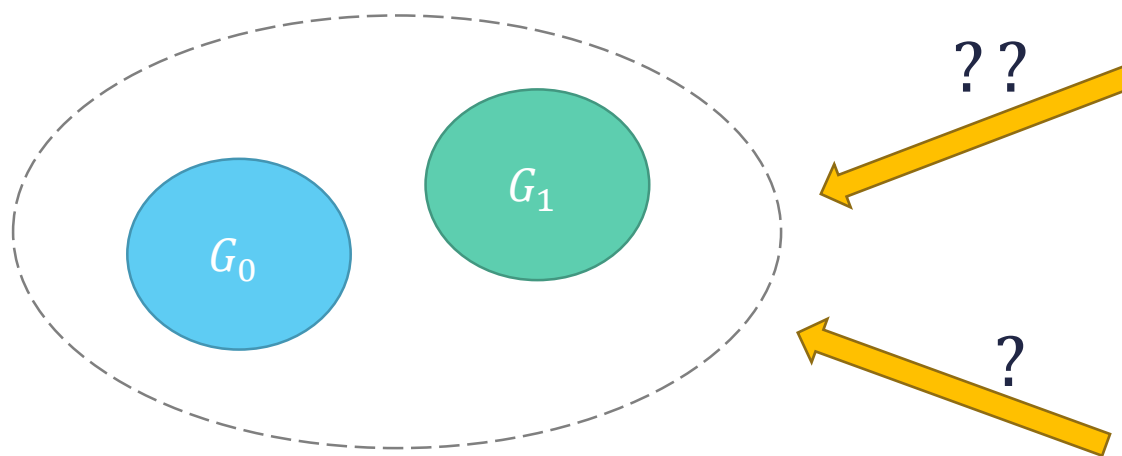
- distinguishing/hypothesis testing
- recovery
- concluding

outline

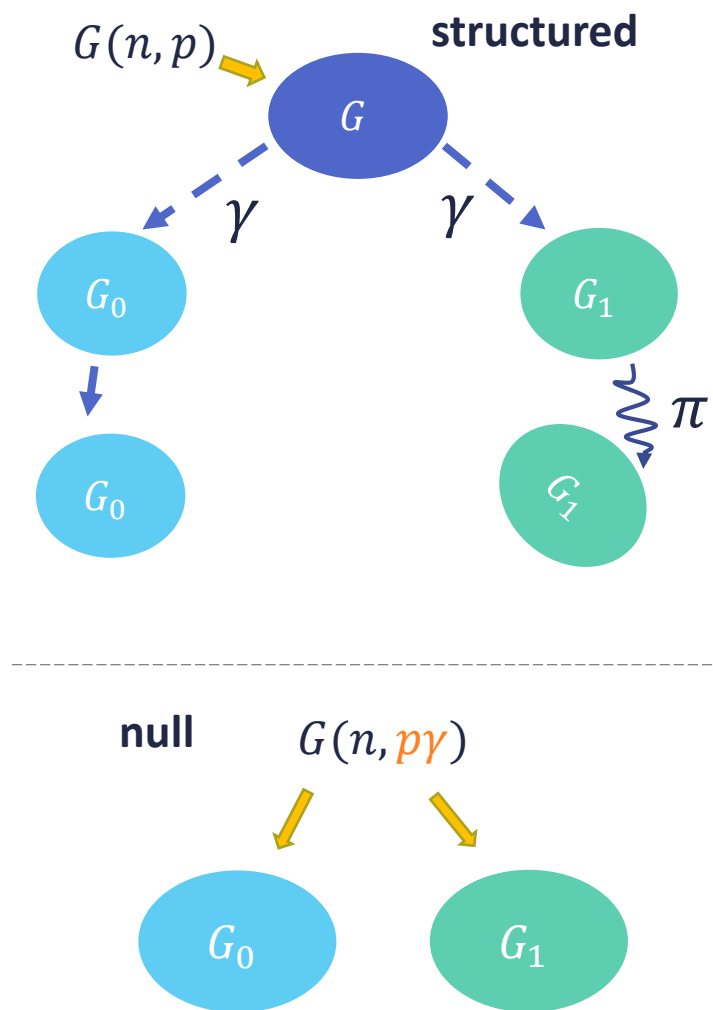
- distinguishing/hypothesis testing
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distinguishing/hypothesis testing

Given G_0, G_1 sampled equally likely from **structured** or **null**,
decide w/prob $1 - o(1)$ from which.

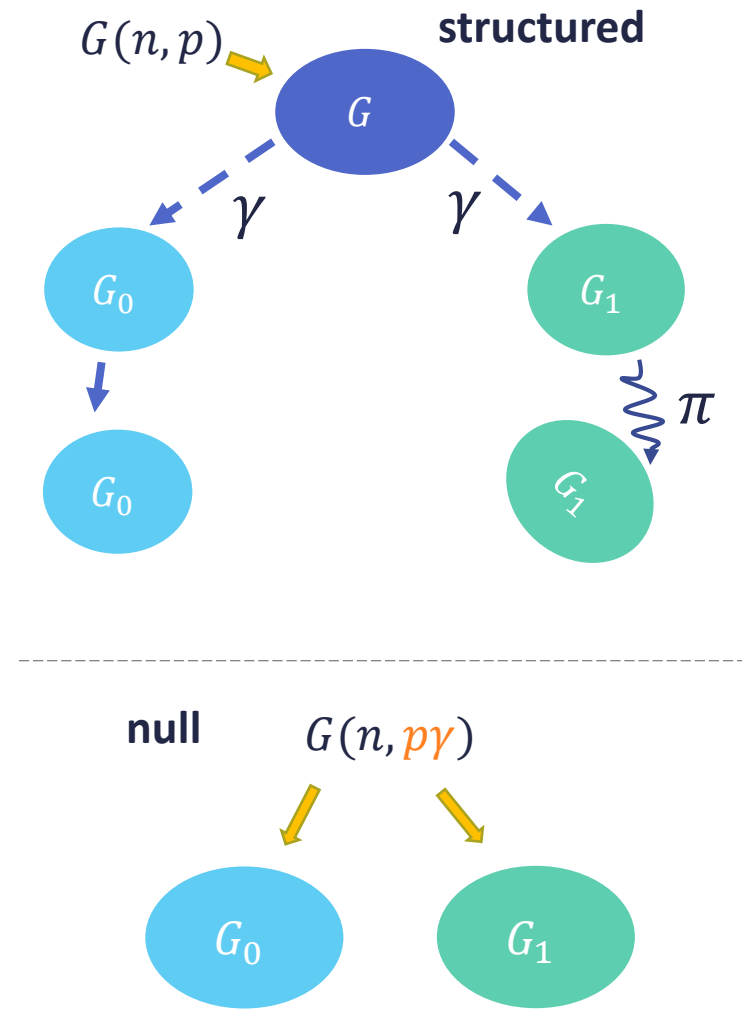
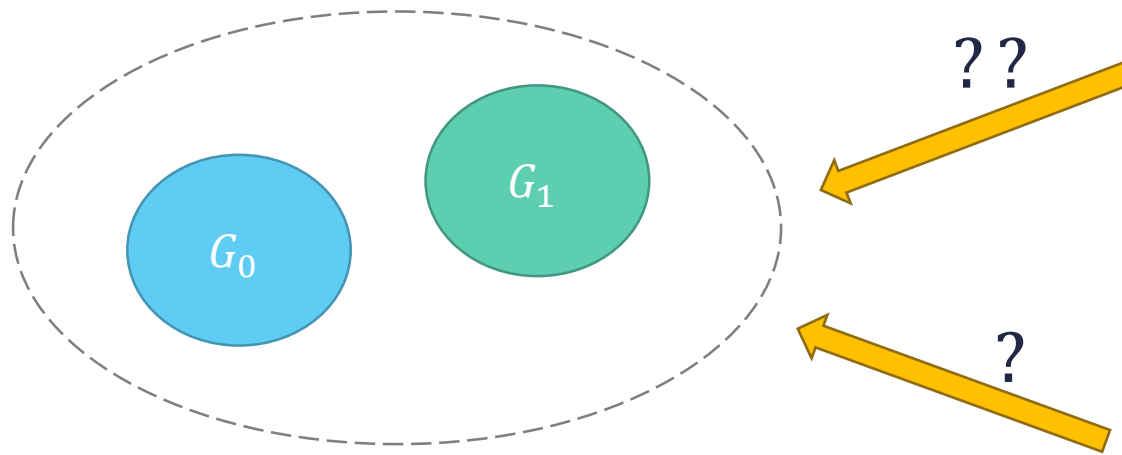


brute force: is there a π with $\geq p\gamma^2 n^2$ matched edges?



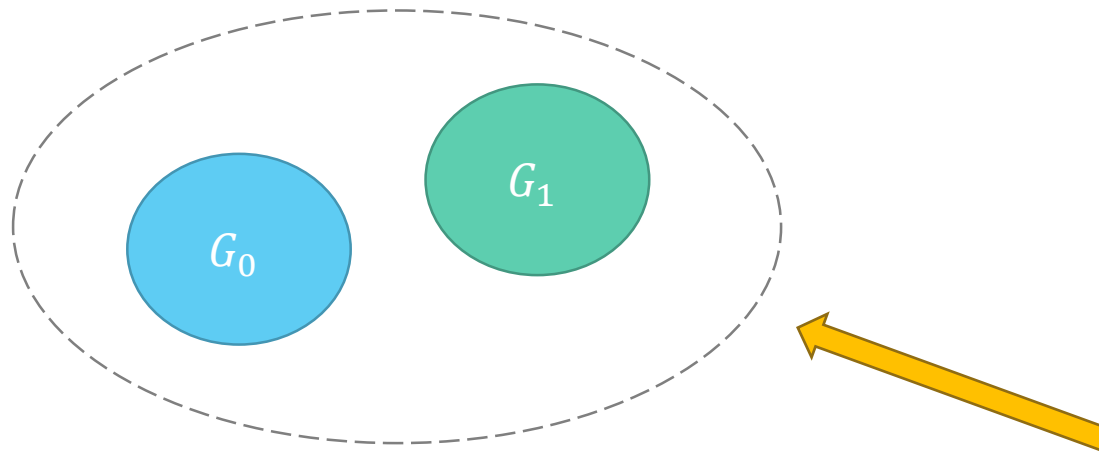
...counting triangles?

$$\text{cor}_{K_3}(G_0, G_1) := (\# K_3 \text{ in } G_0)(\# K_3 \text{ in } G_1).$$



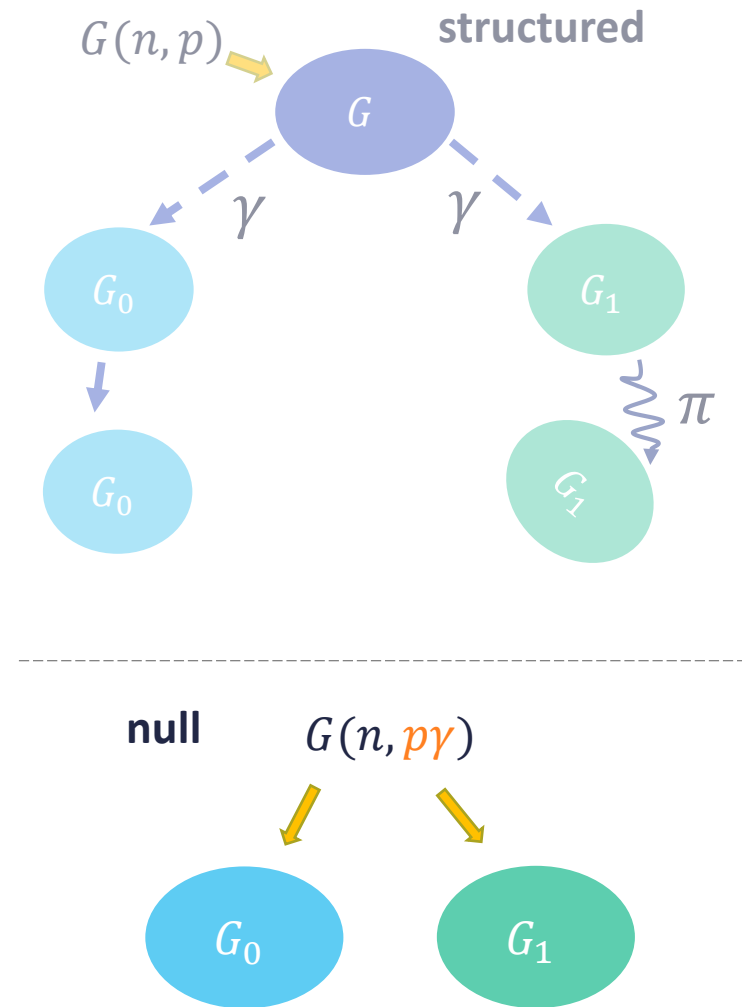
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triangle counts in G_0, G_1 are **independent**

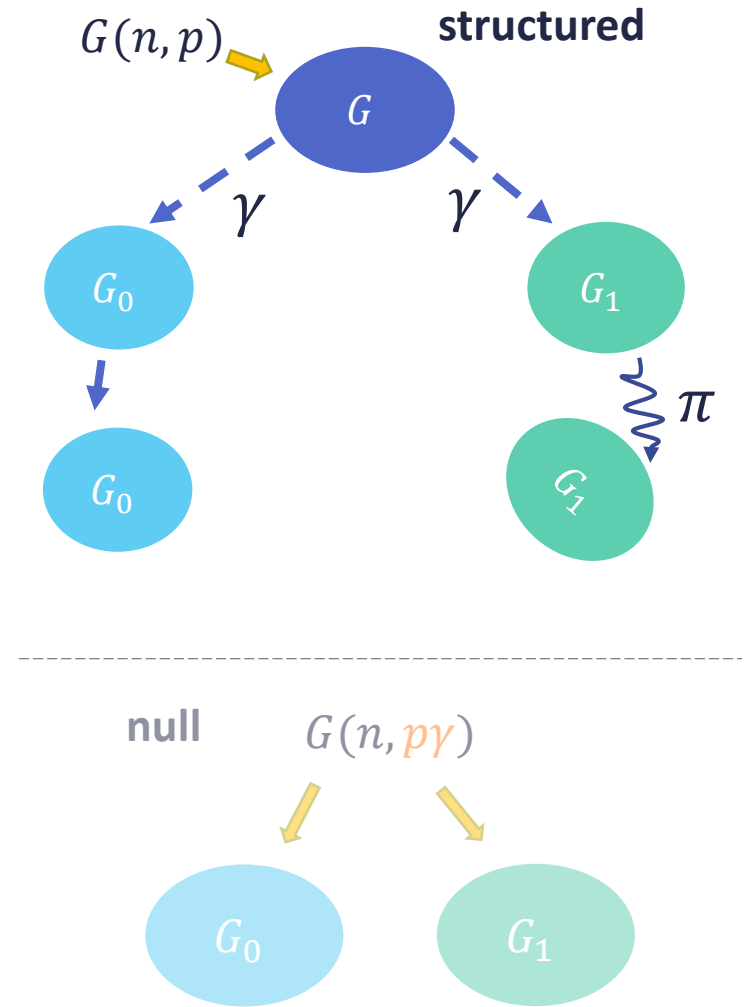
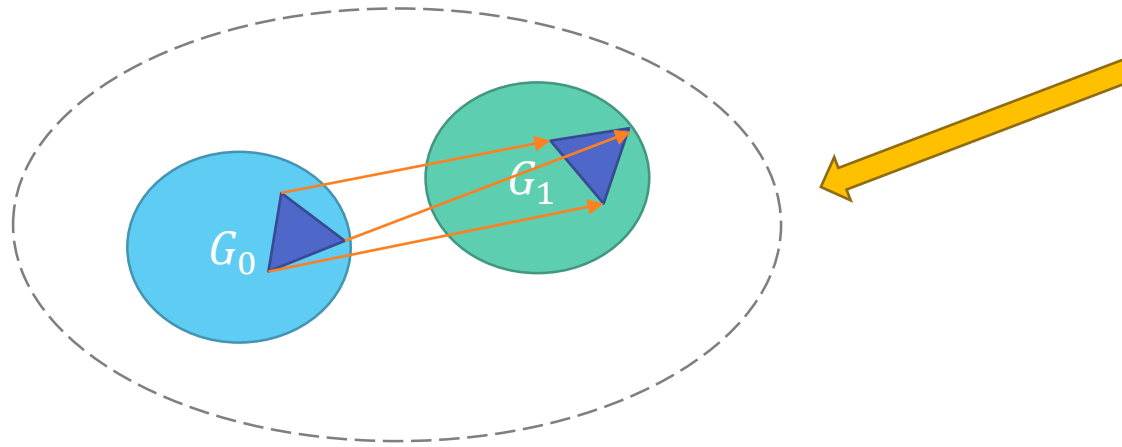
$$\mathbb{E}[\text{cor}_{K_3}(G_0, G_1)] \approx (p\gamma n)^6$$



...counting triangles?

$$\mathbb{E}[cor_{K_3}(G_0, G_1)] \approx (p\gamma n)^6 + (\gamma^2 p n)^3$$

triangle counts in G_0, G_1 are **correlated**



...counting triangles?

structured

$$\mathbb{E}[cor_{K_3}(G_0, G_1)] \approx (p\gamma n)^6 + (\gamma^2 pn)^3$$

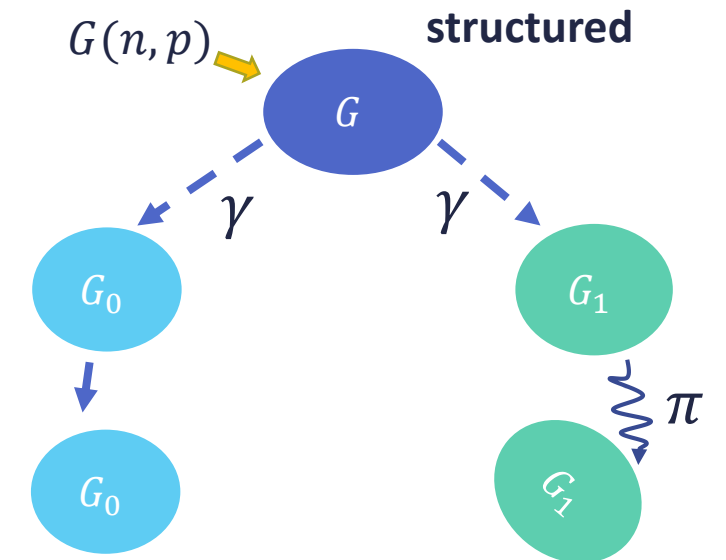
null

$$\mathbb{E}[cor_{K_3}(G_0, G_1)] \approx (p\gamma n)^6$$

Variance?

Optimistically, in null case,

$$\mathbb{V}[cor_{K_3}(G_0, G_1)]^{1/2} \approx (p\gamma n)^3$$



null

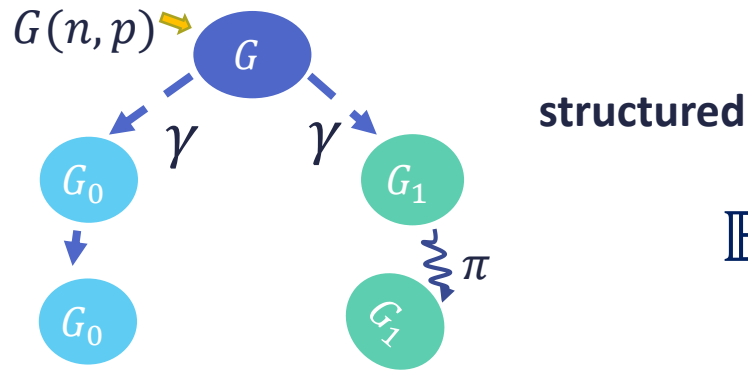
$G(n, p\gamma)$



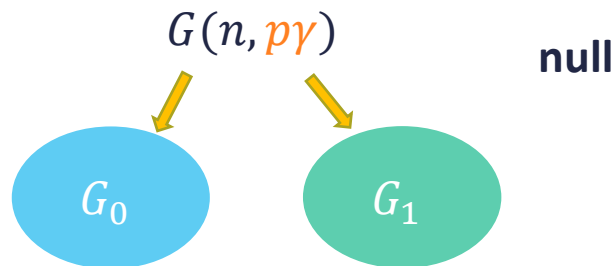
“independent trials”

Suppose we had T “independent trials”:

$$cor_T(G_0, G_1) = \frac{1}{T} \sum_{i=1}^T cor_{K_3}^{(i)}(G_0, G_1)$$



$$\mathbb{E}[cor_T(G_0, G_1)] \approx (p\gamma n)^6 + (\gamma^2 pn)^3$$



$$\mathbb{E}[cor_T(G_0, G_1)] \approx (p\gamma n)^6$$

$$\mathbb{V}[cor_T(G_0, G_1)]^{1/2} \approx \frac{1}{\sqrt{T}} (\gamma^2 pn)^3$$

if $T > 1/\gamma^6$,
 cor_T is a good test

near-independent subgraphs

~~“independent trials”~~

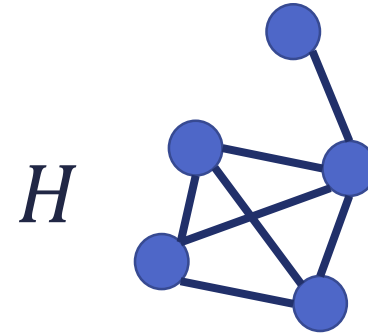
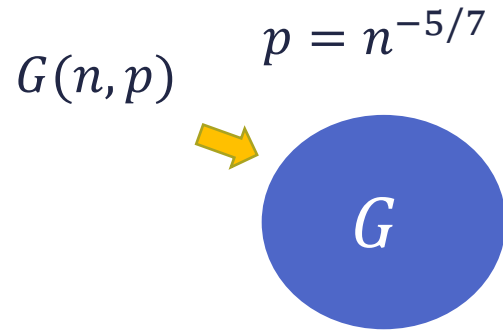
Suppose we had T “independent” *subgraphs*:

H_1, \dots, H_T

$$cor_T(G_0, G_1) = \frac{1}{T} \sum_{i=1}^T cor_{H_i}(G_0, G_1)$$

what properties must H_1, \dots, H_T have to be “independent”?

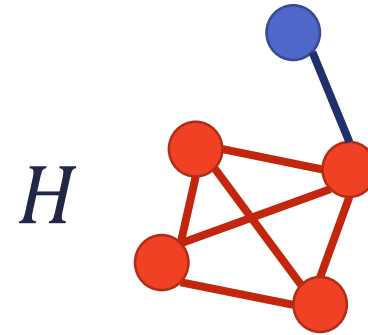
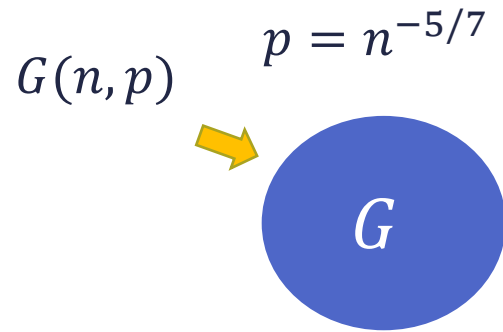
surprisingly delicate (concentration)



How many labeled copies of H in G ?

$$\mathbb{E}[\#_H(G)] = \frac{5!}{|aut(H)|} \cdot \binom{n}{5} \cdot p^7 \approx n^5 p^7 = \Theta(1)$$

surprisingly delicate (concentration)



$\#_H(G)$ does not
concentrate!

How many labeled copies of H in G ?

$$\mathbb{E}[\#_H(G)] = \frac{5!}{|aut(H)|} \cdot \binom{n}{5} \cdot p^7 \approx n^5 p^7 = \Theta(1)$$

How many labeled copies of K_4 in G ?

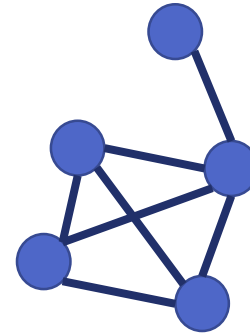
$$\mathbb{E}[\#_{K_4}(G)] = \frac{4!}{|aut(K_4)|} \cdot \binom{n}{4} \cdot p^6 \approx n^4 p^6 = \Theta(n^{-2/7})$$

variance of subgraph counts

$G(n, p)$



H



Lemma

For a constant-sized subgraph H ,

$$\mathbb{V}[\#_H(G)] = \Theta(1) \cdot \frac{\mathbb{E}[\#_H(G)]^2}{\min_{J \subset H} \mathbb{E}[\#_J(G)]}$$

subgraph of H with fewest expected appearances

strict balance

H is *strictly balanced* if all its strict subgraphs have edge density $< \frac{|E(H)|}{|V(H)|}$.

↳ if $\mathbb{E}[\#_H(G)] \approx n^{|V(H)|} p^{|E(H)|} = \Theta(1)$,

then $\mathbb{E}[\#_J(G)] = \omega(1)$ for any $J \subset H$.

Lemma

For a constant-sized subgraph H ,

$$\mathbb{V}[\#_H(G)] = \Theta(1) \cdot \frac{\mathbb{E}[\#_H(G)]^2}{\min_{J \subset H} \mathbb{E}[\#_J(G)]} = o(1) \cdot \mathbb{E}[\#_H(G)]$$

concentration AND independence

If H_1, \dots, H_T are non-isomorphic strictly balanced graphs with $\mathbb{E}[\#_{H_i}(G)] = \Theta(1)$,

their counts concentrate

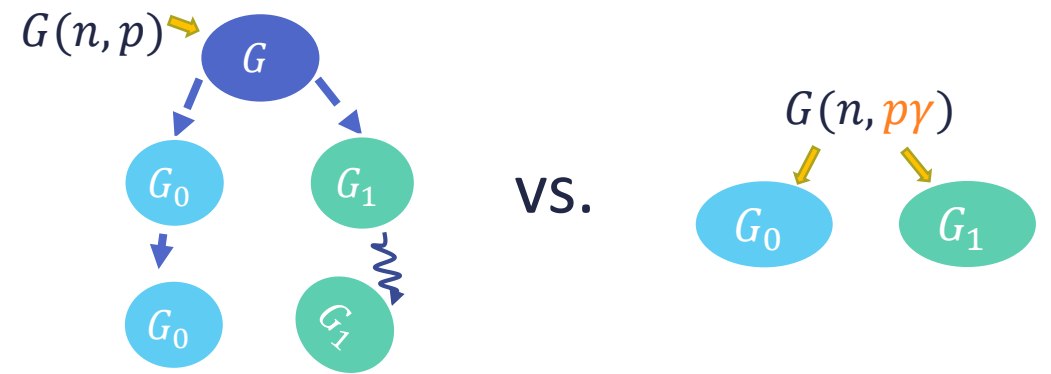
$$\forall i \in [T], \quad \mathbb{V}[\#_{H_i}(G)] = o(1) \cdot \mathbb{E}[\#_{H_i}(G)]$$

their counts are asymptotically independent

$$\forall i \neq j \in [T],$$

$$\mathbb{E}[\#_{H_i}(G) \cdot \#_{H_j}(G)] = (1 + o(1)) \cdot \mathbb{E}[\#_{H_i}(G)] \cdot \mathbb{E}[\#_{H_j}(G)]$$

distinguishing algorithm

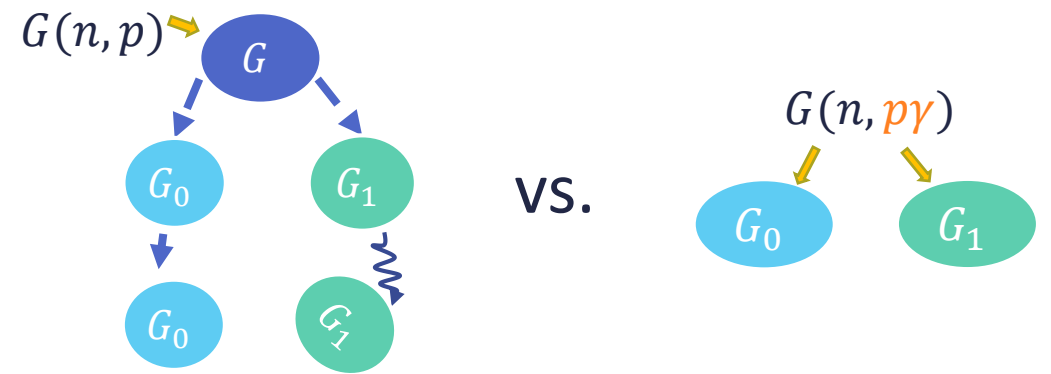


For $\nu = \frac{1}{\text{poly}(\gamma)}$, design a “test set” H_1, \dots, H_T

of $T = \nu^{\Omega(e)}$ **strictly balanced** graphs w/ ν vertices & e edges.

set $n^\nu (p\gamma)^e \approx 1$

distinguishing algorithm



For $v = \frac{1}{\text{poly}(\gamma)}$, design a “test set” H_1, \dots, H_T

of $T = v^{\Omega(e)}$ **strictly balanced** graphs w/ v vertices & e edges.

set $n^v(p\gamma)^e \approx 1$

compute

$$\text{cor}_T(G_0, G_1) = \frac{1}{T} \sum_{i=1}^T \text{cor}_{H_i}(G_0, G_1)$$

$\geq \theta$

structured

$< \theta$

null

$$\mathbb{E}[\text{cor}_T(G_0, G_1)] = \begin{cases} n^{2v}(\gamma p)^{2e} + n^v(\gamma^2 p)^e & \text{structured} \\ n^{2v}(\gamma p)^{2e} & \text{null} \end{cases}$$

TODO: variance in structured case.

$$\mathbb{V}[\text{cor}_T(G_0, G_1)] = \frac{1}{\sqrt{T}} n^v(\gamma p)^e < n^v(\gamma^2 p)^e \quad \text{null}$$

outline

- distinguishing/hypothesis testing
- recovery
- concluding

outline

- distinguishing/hypothesis testing
- test graphs
- recovery
- concluding

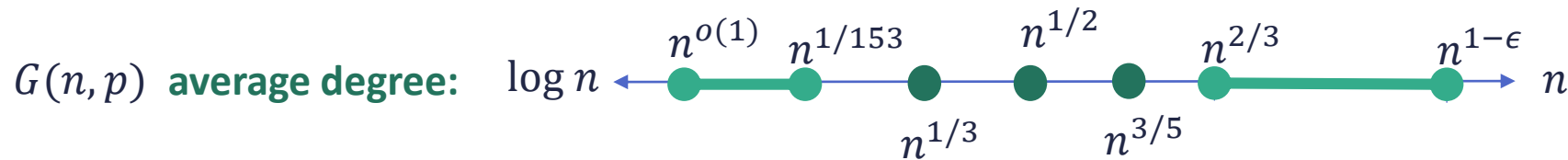
designing a “test set”

For $v = \frac{1}{\text{poly}(\gamma)}$, design a “test set” H_1, \dots, H_T

of $T = v^{\Omega(e)}$ **strictly balanced** graphs w/ v vertices & e edges.

$$\text{set } n^v (p\gamma)^e \approx 1$$

remember?



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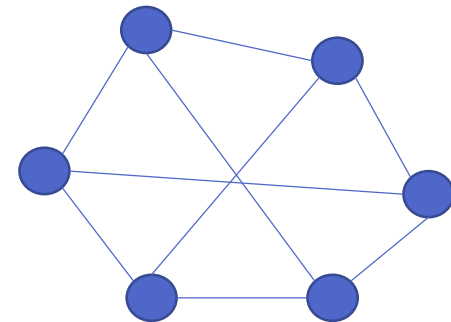
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claim: connected d -regular graphs are **strictly balanced**.

proof: in any strict subgraph, average degree $< d$.



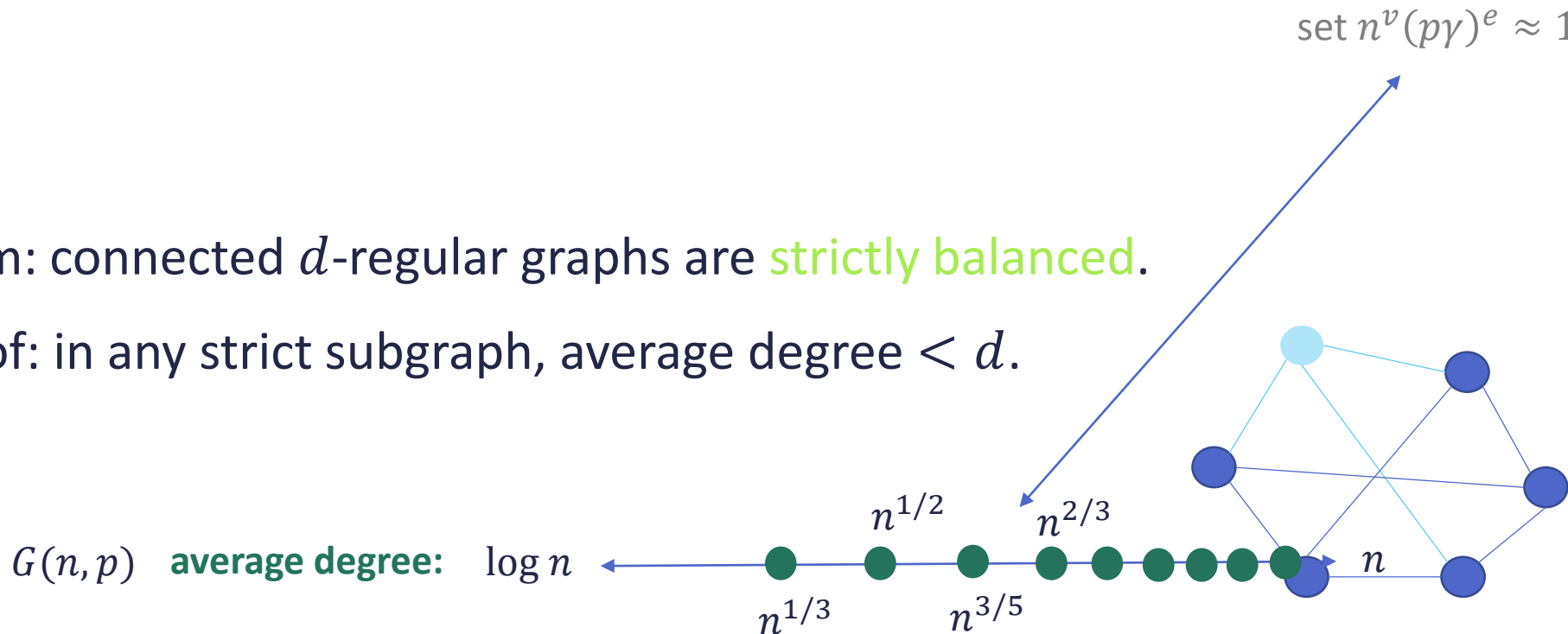
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“test set” for non-integer degrees

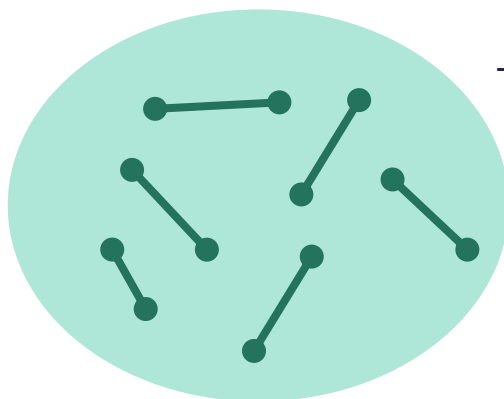
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set $n^v(p\gamma)^e \approx 1$

what if we want $2 \cdot \frac{e}{v} = \lambda \cdot (d + 1) + (1 - \lambda) \cdot d$?

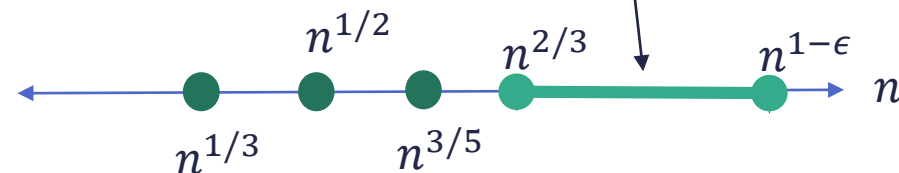
d -regular random graph on v vertices



+ random matching on λv vertices

strict balance? expansion.

$G(n, p)$ average degree: $\log n$



“test set” for non-integer degrees

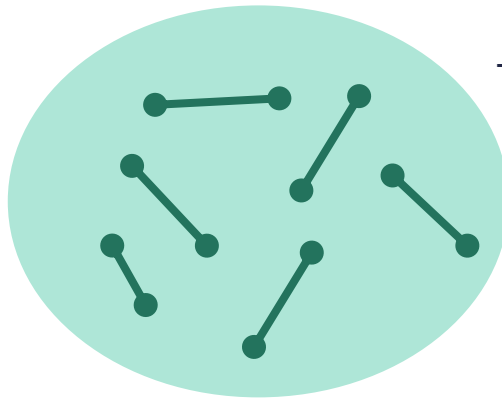
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what if we want $2 \cdot \frac{e}{v} = \lambda \cdot (d + 1) + (1 - \lambda) \cdot d$?

d -regular random
graph on v vertices



+ random matching
on λv vertices

strict balance? expansion.

$d < 3$?

2-regular graphs don't expand.

“test set” for non-integer degrees < 3

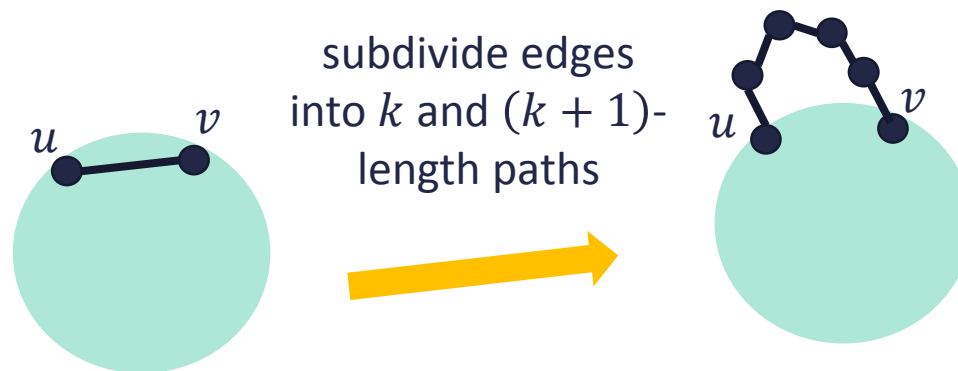
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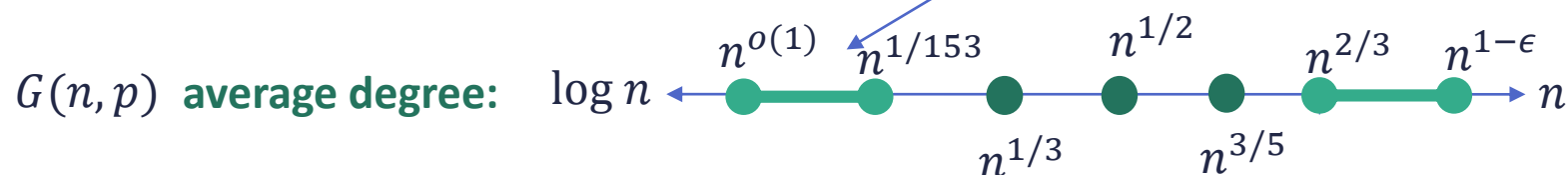
$$\text{set } n^v (p\gamma)^e \approx 1$$

what if we want $2 \cdot \frac{e}{v} = \lambda \cdot 3 + (1 - \lambda) \cdot 2$?

3-regular random graph on λv vertices



strict balance? expansion.



designing a “test set”

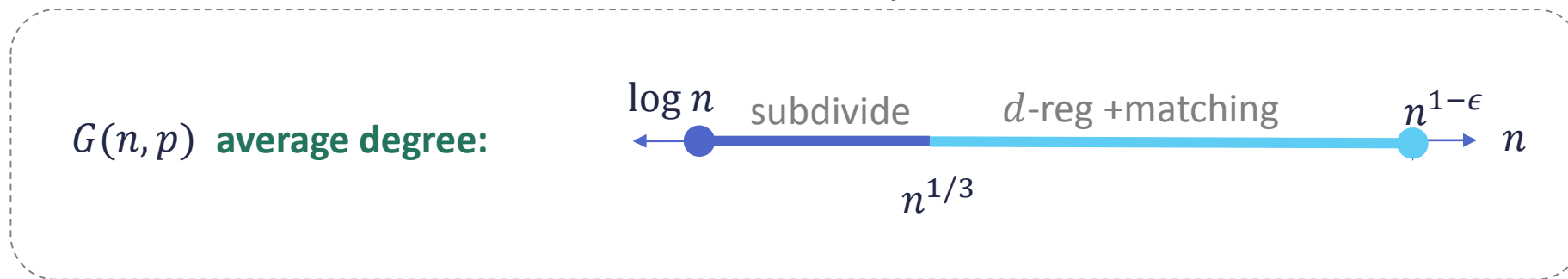
For $v = \frac{1}{\text{poly}(\gamma)}$, design a “test set” H_1, \dots, H_T

of $T = v^{\Omega(e)}$ **strictly balanced** graphs w/ v vertices & e edges.

set $n^v(p\gamma)^e \approx 1$

+ more conditions (for recovery)

Conjecture: our construction achieves all $\frac{e}{v}$



outline

- distinguishing/hypothesis testing
- test graphs
- recovery
- concluding

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distinguishing \neq recovery

distinguishing: counting subgraphs



ambiguity in matching; how to conclude $\pi(u) = v$?

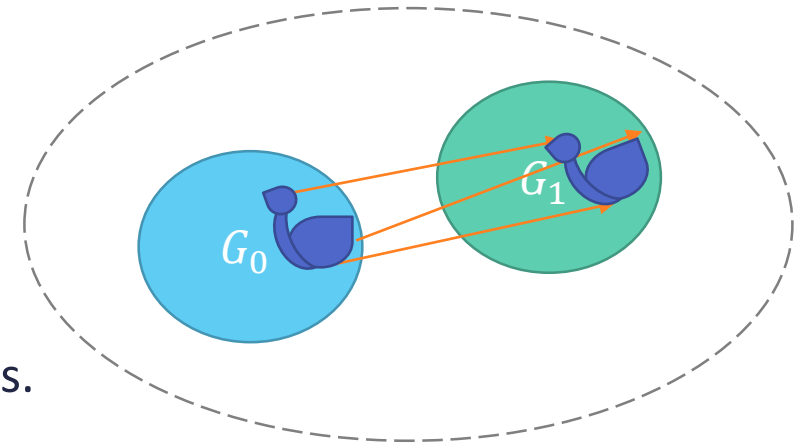
distinguishing: subgraphs on $\frac{1}{\text{poly}(\gamma)} = O(1)$ vertices, each appearing $O(1)$ times




only $O(1)$ vertices participate in subgraphs from our test set.

the “black swan” approach

identify rare subgraphs appearing in both graphs, and match vertices.



expected
number of  that survive
subsampling

choose test set H_1, \dots, H_T so that $(\gamma^2 p)^e n^v \gg (\gamma p)^{2e} n^{2v}$  expected number of
unrelated pairs of



if we see H_i in both graphs, it is most likely because of correlation.

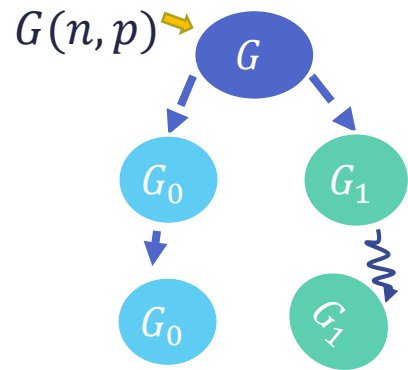
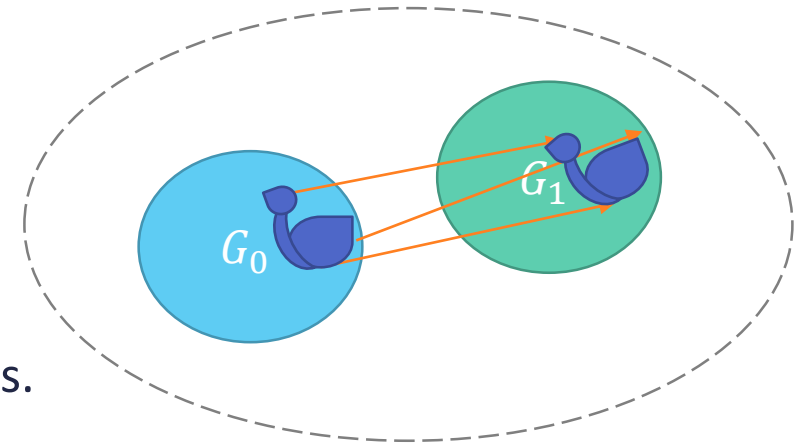
choose large test set H_1, \dots, H_T with $v = O(\log n)$ vertices




$\Omega(n)$ vertices participate in subgraphs from our test set.

the “black swan” approach

identify rare subgraphs appearing in both graphs, and match vertices.



Claim: there is at most one copy of each  in G with high probability

Claim: $\Omega(n)$ vertices in $G_0 \cap G_1$, appear in a surviving subsampled  with high probability

proofs: second moment method

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why subgraph counts/statistics?

emerging intuition/conjectures: $\text{SoS} \equiv_{avg} \text{low-degree polynomials}$

the sum-of-squares (SoS) semidefinite program is at most as powerful as “low-degree” statistics for average-case problems.

known to hold for: planted clique [Barak-Hopkins-Kelner-Kothari-Moitra-Potechin’16]

CSP refutation [Grigoriev’01, Schoenebeck’08, Kothari-Mori-O’Donnell-Witmer’17]

tensor PCA [Hopkins-Kothari-Potechin-Raghavendra-S-Steurer’17]

also known: SoS is at most as powerful as “low-degree” spectral algorithms for average-case problems [Hopkins-Kothari-Potechin-Raghavendra-S-Steurer’17]

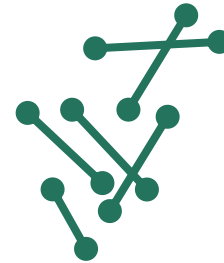
does SoS *know* about the black swans?

does the natural SoS relaxation recover π ?

$$\max_{\pi} \langle A_{G_0}, \pi(A_{G_1}) \rangle$$

SoS relaxation

cares
about



or



?

can ask similar questions about other low-degree functions,
e.g. non-backtracking random walk matrix.

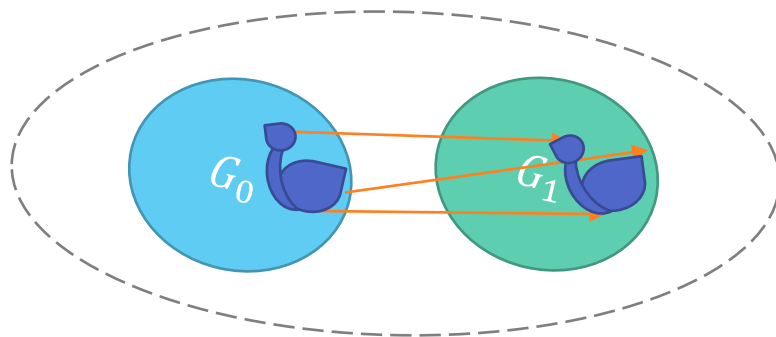
more questions

- recovery in polynomial time?

SoS? or, many variations on our theme are possible.

- all information-theoretically possible $p \in \left[\frac{\log n}{n}, O(1) \right]$?

- practical heuristics?



Thank you!