

A POLYNOMIAL-TIME APPROXIMATION ALGORITHM FOR ALL-TERMINAL NETWORK RELIABILITY

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Queen Mary Algorithm Day, Jul 17, 2018

Random sampling strikes back

The complexity of computing quantities

Complexity class **#P** by Valiant (1979):

a counting analogue of **NP**.

Evaluation of probabilities;

Partition functions in statistical physics;

Counting discrete structures ...



The complexity of approximate counting

What about (multiplicatively) approximating **#P**-complete problems?

- at most **NP**-hard (Valiant and Vazirani, 1986);
- any polynomial approximation can be amplified into an ϵ -approximation with polynomial overhead.

Efficient approximation algorithms do exist! Famous examples include

- the volume of a convex body
(Dyer, Frieze, and Kannan, 1991);
- the partition function of ferromagnetic Ising models
(Jerrum and Sinclair, 1993);
- the permanent of a non-negative matrix
(Jerrum, Sinclair, and Vigoda, 2004).

There are still many open problems in approximate counting!

NETWORK RELIABILITY



Network reliability

Given a undirected graph (a.k.a. network) $G = (V, E)$, define a random subgraph $G(p)$ by **removing** each edge independently with probability p .

(ALL-TERMINAL) RELIABILITY is the probability such that $G(p)$ is connected.

One may ask the probability of other properties of $G(p)$, such as whether two distinct vertices s and t are connected (**s-t RELIABILITY**), or whether $G(p)$ is acyclic (**counting weighted forests**), etc.

Network reliability

(ALL-TERMINAL) RELIABILITY: The probability that $G(p)$ is connected.

In other words, we want to compute

$$Z_{\text{rel}}(G, p) := \sum_{R \subseteq E: (V, R) \text{ is connected}} p^{|E \setminus R|} (1-p)^{|R|}.$$

For example:

$$Z_{\text{rel}}(\text{---} \circ \text{---} \circ \text{---} \circ \text{---} \circ \text{---}, p) = \text{---} \circ \text{---} \circ \text{---} \circ \text{---} \circ \text{---} = (1-p)^{n-1};$$

$$\begin{aligned} Z_{\text{rel}}(\square, p) &= \square + \square + \square + \square + \square \\ &= (1-p)^4 + 4p(1-p)^3; \end{aligned}$$

$$Z_{\text{rel}}(G, 1/2) = \frac{|\{R \subseteq E : (V, R) \text{ is connected}\}|}{2^{|E|}}.$$

Computational complexity of reliability

Directed and undirected **s-t RELIABILITY** (and a few other variants) are featured in the original list of 13 **#P**-complete problems by Valiant (1979).

Exact evaluation of **ALL-TERMINAL RELIABILITY** is shown to be **#P**-complete by Jerrum (1981), and independently Provan and Ball (1983).

What about approximation? Open since 80s.

Karger (1999) has given a famous FPRAS for **UNRELIABILITY** (namely $1 - Z_{rel}$). However, approximating $1 - Z_{rel}$ does not yield a good approximation for Z_{rel} when Z_{rel} is exponentially small.

The Tutte polynomial

For a **connected** undirected graph $G = (V, E)$,

$$Z_{\text{Tutte}}(G; x, y) := \sum_{R \subseteq E} (x-1)^{\kappa(R)-1} (y-1)^{\kappa(R)+|R|-|V|},$$

where $\kappa(R)$ is the number of connected components of (V, R) .

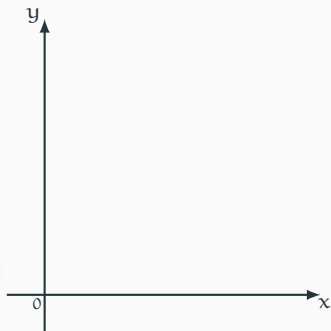
$(1, 1)$: counting spanning trees;

$(1, 1/p)$: **network reliability**;

$(x, 1)$: counting weighted forests;

$(x-1)(y-1) = 2$:
ferromagnetic Ising model;

and many more ...



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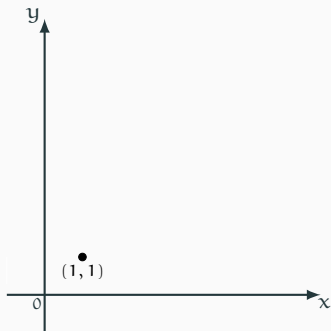
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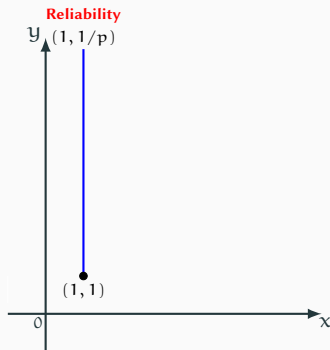
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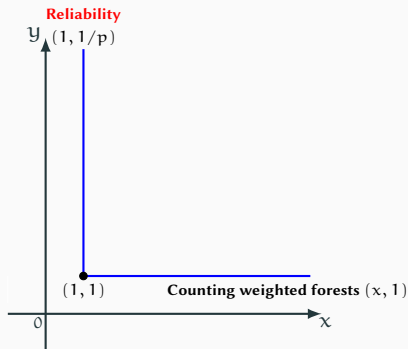
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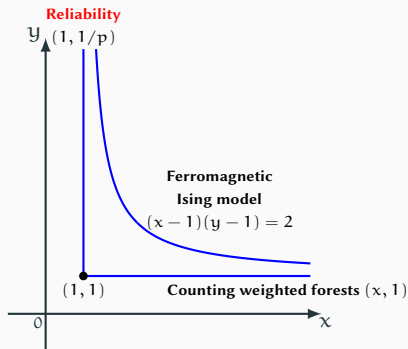
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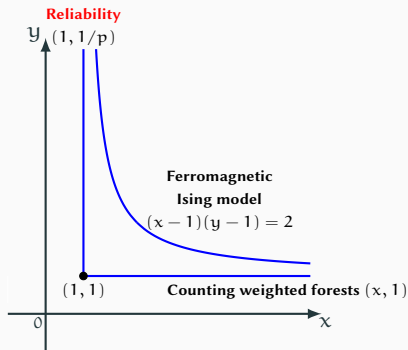
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Main result

Let $m := |E|$ and $n := |V|$.

Theorem (G. and Jerrum, 2018)

There is a randomised algorithm approximating Z_{rel} within multiplicative factor $(1 \pm \epsilon)$, with expected running time $O(\epsilon^{-2}(1-p)^{-3}m^2n^3)$.

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There is an exact sampler to draw (edge-weighted) connected subgraphs with expected running time $O((1-p)^{-1}m^2n)$.

Spoiler: sampling can be done in $O(mn)$ time and approximate counting in $O(mn^2 \log n)$ time (G. and He, 2018+).

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NATURAL ATTEMPTS

(AND WHY THEY DO NOT SUCCEED)



Naive Monte Carlo

A natural unbiased estimator \tilde{Z} of Z_{rel} :

1. Draw k independent subgraphs $(R_i)_{i \in [k]}$ of $G(p)$.
2. Let

$$\tilde{Z} := \frac{1}{k} \sum_{i \in [k]} \mathbb{1}_{\text{conn}}(R_i),$$

where $\mathbb{1}_{\text{conn}}(R)$ is the indicator variable of (V, R) being connected.

It is easy to see that $\mathbb{E} \tilde{Z} = Z_{\text{rel}}$.

However, if Z_{rel} is exponentially small (e.g. $Z_{\text{rel}}(P_n, p) = (1-p)^{n-1}$), then we will almost never see a connected R_i .

In that case, the variance of $\mathbb{1}_{\text{conn}}(R)$ is exponentially large, and k has to be exponentially large to yield a good approximation.

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Nonetheless, naive Monte Carlo (NMC) is the basic building block of the FPRAS by [Karger \(1999\)](#) for **UNRELIABILITY** (namely $1 - Z_{\text{rel}}$).

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Reducing counting to sampling

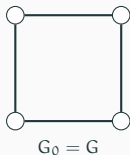
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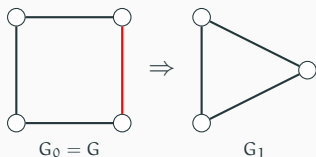
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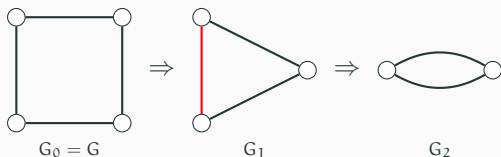
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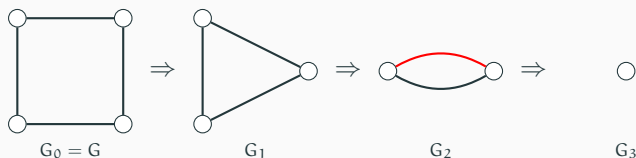
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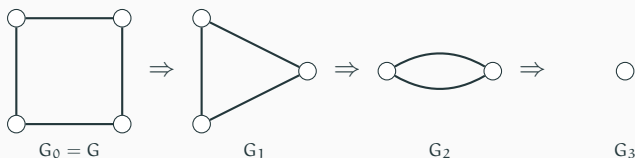
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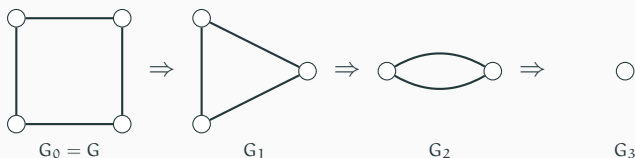
Rewrite

$$Z_{\text{rel}}(G) = \frac{Z_{\text{rel}}(G_0)}{Z_{\text{rel}}(G_1)} \cdot \frac{Z_{\text{rel}}(G_1)}{Z_{\text{rel}}(G_2)} \cdot \frac{Z_{\text{rel}}(G_2)}{Z_{\text{rel}}(G_3)} \cdot Z_{\text{rel}}(G_3).$$

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To estimate $\frac{Z_{\text{rel}}(G_i)}{Z_{\text{rel}}(G_{i+1})}$, draw $C \sim \pi_{G_{i+1}}(\cdot)$ and let

$$C' := \begin{cases} C & \text{with prob. } p; \\ C \cup \{e\} & \text{otherwise,} \end{cases} \quad \text{and} \quad X := \mathbb{1}_{\text{conn}, G_i}(C').$$

Then $\mathbb{E}X = \frac{Z_{\text{rel}}(G_i)}{Z_{\text{rel}}(G_{i+1})}$ and its variance is bounded by a constant.

Markov chain Monte Carlo

Markov chains is the “off the shelf” approach to sampling from complicated distributions.

There is a natural Markov chain converging to $\pi_G(\cdot)$:

1. Let $C_0 = E$.

2. Given C_t , randomly pick an edge $e \in E$.

If $C_t \setminus \{e\}$ is disconnected then $C_{t+1} = C_t$. Otherwise,

$$C_{t+1} = \begin{cases} C_t \cup \{e\} & \text{with prob. } 1 - p; \\ C_t \setminus \{e\} & \text{with prob. } p. \end{cases}$$

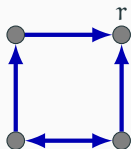
Unfortunately, no polynomial upper bound (nor exponential lower bound) is known about its mixing time (rate of convergence).

A SURPRISING EQUIVALENCE

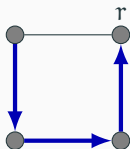
(AND AN ALTERNATIVE WAY TO SAMPLING)

Reachability

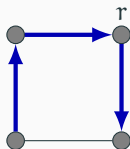
We say a directed graph D with root r is *root-connected* if all vertices can reach r .



Root-connected



Root-connected



Not root-connected!

REACHABILITY: in a directed graph $D = (V, A)$ with root r , what's the probability that $D(p)$ is root-connected?

$$Z_{\text{reach}}(D, p) := \sum_{R \subseteq A: (V, R) \text{ is root-connected}} p^{|A \setminus R|} (1 - p)^{|R|}.$$

A surprising equivalence

Ball (1980) showed that for any undirected graph $G = (V, E)$,

$$Z_{\text{rel}}(G, p) = Z_{\text{reach}}(\vec{G}, p),$$

where \vec{G} is the directed graph obtained by replacing every $e \in E$ with a pair of anti-parallel arcs. (Called **bi-directed**).



Thus we just need to approximate **REACHABILITY** in bi-directed graphs.

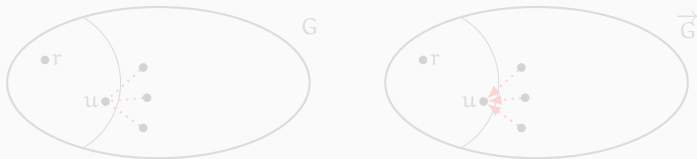
A coupling proof

We have an alternative coupling proof of Ball's equivalence:

There is a coupling \mathcal{C} under which

$$G(p) \text{ is connected} \Leftrightarrow \vec{G}(p) \text{ is root-connected.}$$

Explore G and \vec{G} like a BFS, starting from r . Reveal $\vec{G}(p)$ and $G(p)$ as the process proceeds. Couple the arc going towards the current vertex in $\vec{G}(p)$ with the corresponding edge in $G(p)$.



When both exploration processes end, the sets of vertices that can reach r are exactly the same.

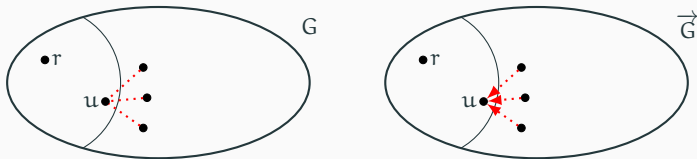
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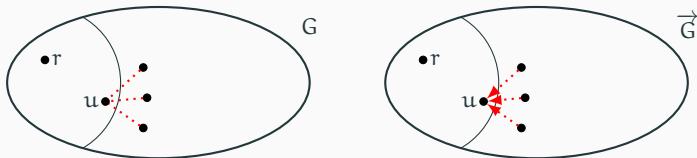
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Sample in at least two ways

Goal: sample uniform (or edge-weighted) root-connected subgraphs.

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However, another hope is to do (exact) sampling in expected polynomial-time, based on rejections.

Cluster-popping

Goal: sample uniform (or edge-weighted) root-connected subgraphs.

Gorodezky and Pak (2014) proposed the “cluster-popping” algorithm:
(Cluster: a subset of vertices not including r and with no arc going out.)

1. Let R be a subset of arcs by choosing each arc e with probability $1 - p$ independently.
2. **While** there is at least one cluster in (V, R) :
 - Let C_1, \dots, C_k be all **minimal** clusters in (V, R) , and $C = \bigcup_{i=1}^k C_i$.
 - Re-randomize all arcs whose heads are in C to get a new R .

Gorodezky and Pak (2014) showed that this algorithm draws from the correct distribution, and they also conjectured that cluster-popping runs in expected polynomial time in bi-directed graphs.

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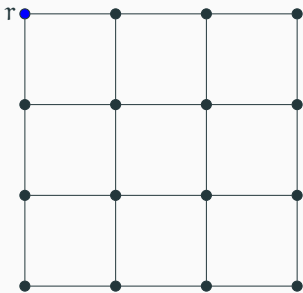
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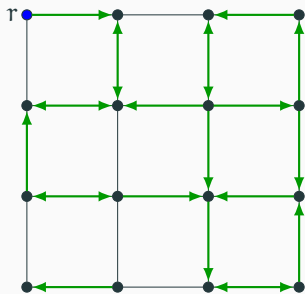
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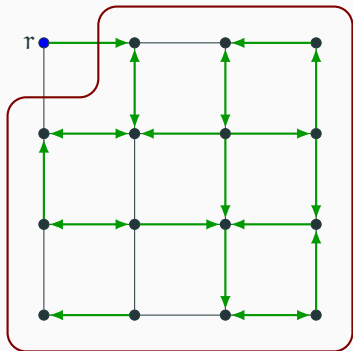
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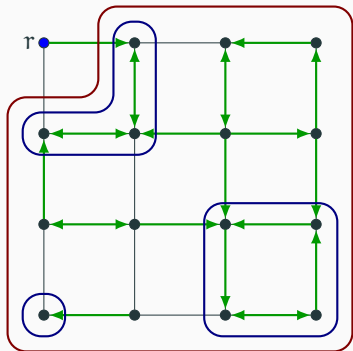
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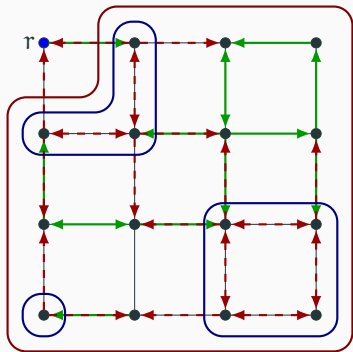
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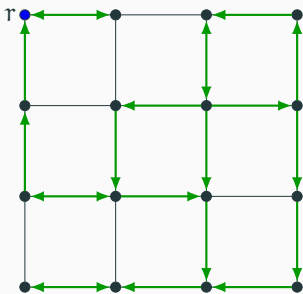
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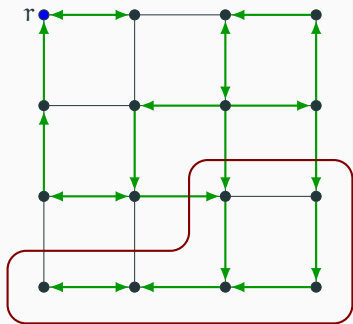
An example run

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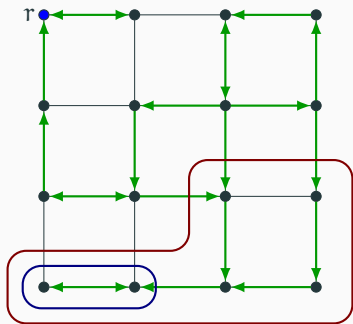
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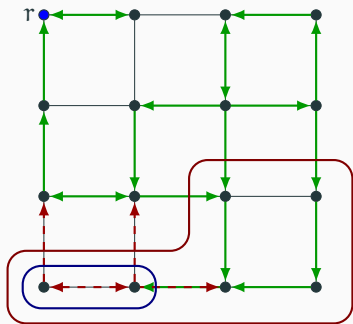
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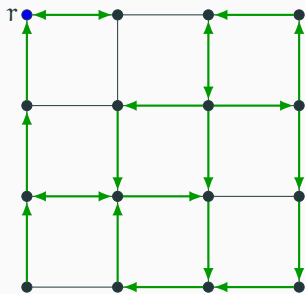
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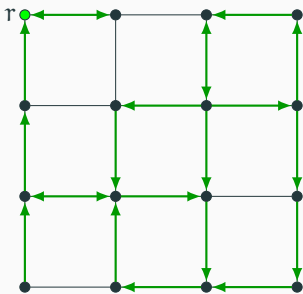
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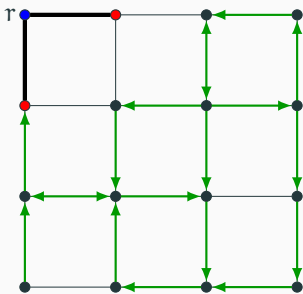
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Mapping back to connected subgraph.
 (Exploration order: left to right, bottom to top)

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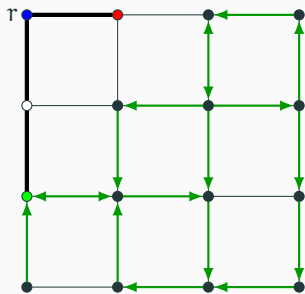


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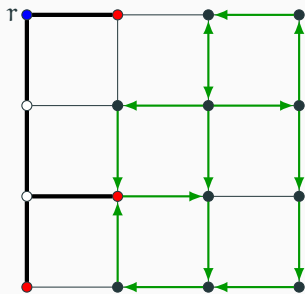


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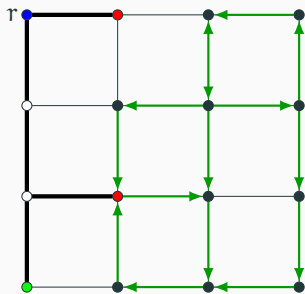


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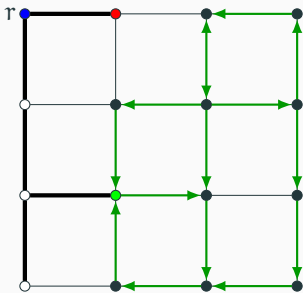


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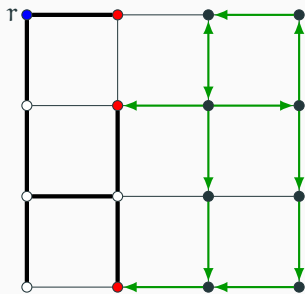


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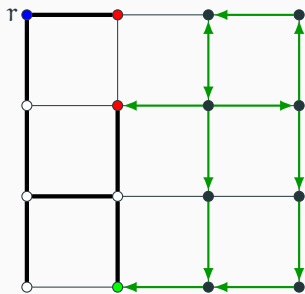


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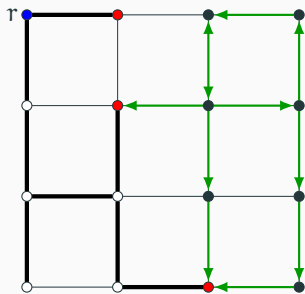
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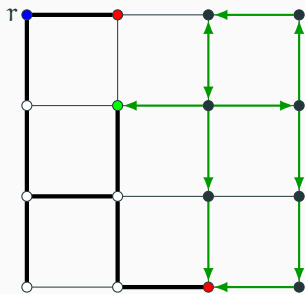
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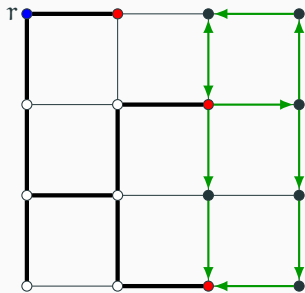
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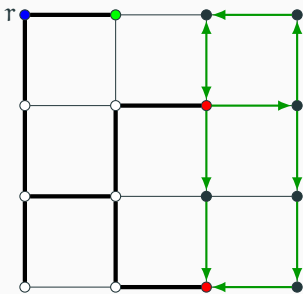
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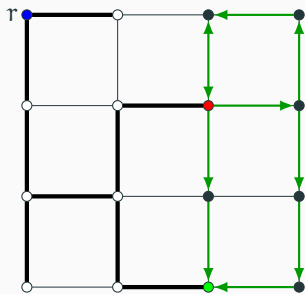
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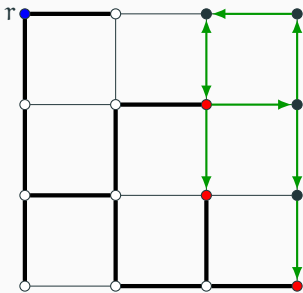
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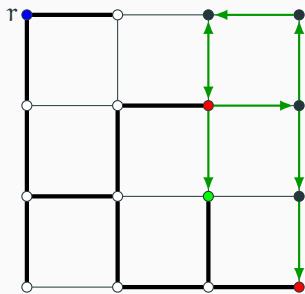
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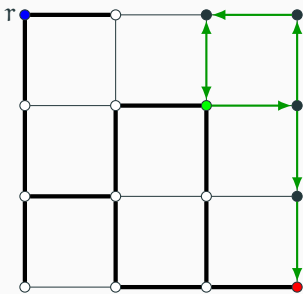
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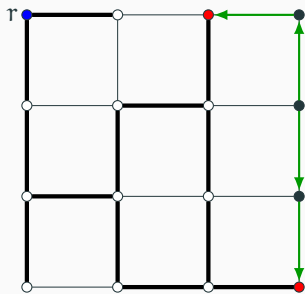
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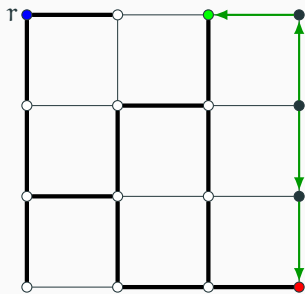
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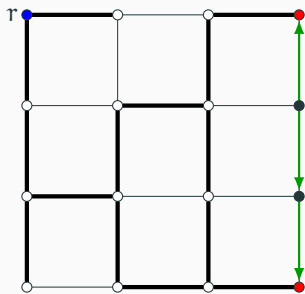
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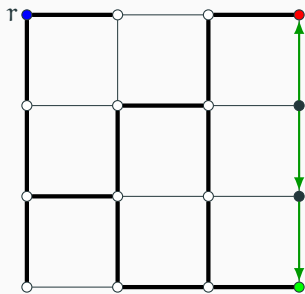
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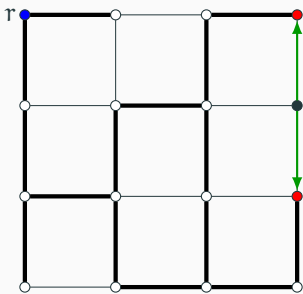
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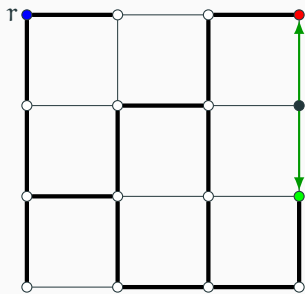
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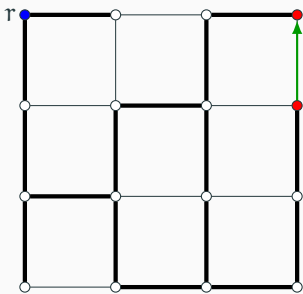
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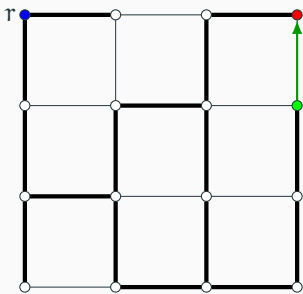
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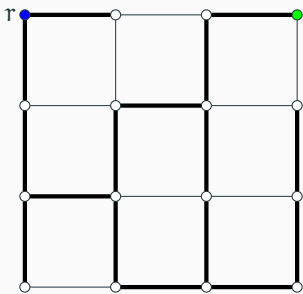


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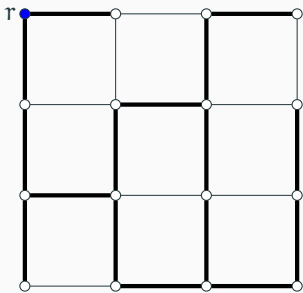
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PARTIAL REJECTION SAMPLING

(A MORE GENERAL PERSPECTIVE FOR CLUSTER-POPPING)

Partial rejection sampling

Cluster-popping is a special case of **PARTIAL REJECTION SAMPLING** framework (G., Jerrum, and Liu, 2017).

The goal is to sample from a product distribution, conditioned on a number of “**bad**” events not happening.

Rejection sampling throws away all variables.

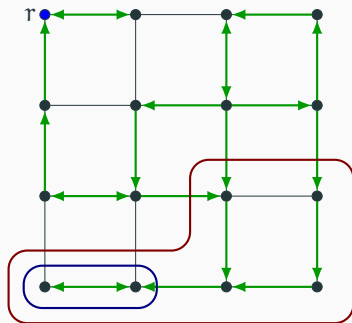
Instead, we want to recycle some randomness while resampling the “**bad**” events (and hopefully not too much more).

Partial rejection sampling

Cluster-popping under partial rejection sampling:

Arcs are variables.

Minimal clusters are “bad” events.

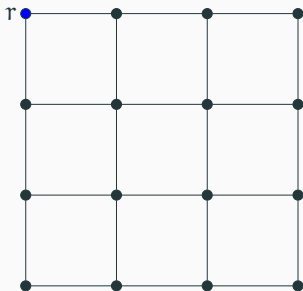


There can be exponentially many bad events.

Extremal instances

An instance is called **extremal** (in the sense of [Shearer \(1985\)](#) regarding non-uniform Lovász Local Lemma):

if any two “bad” events A_i and A_j are either **independent** or **disjoint**.

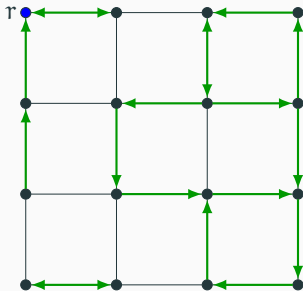


If the instance is **extremal**, then eliminating precisely the “bad” events in each iteration yields the correct distribution once the process halts ([GJL'17](#))!

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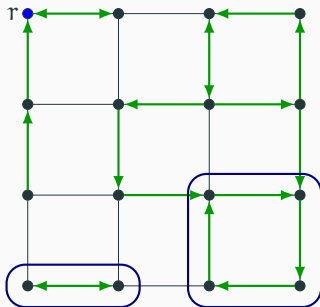


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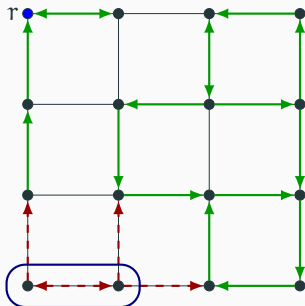


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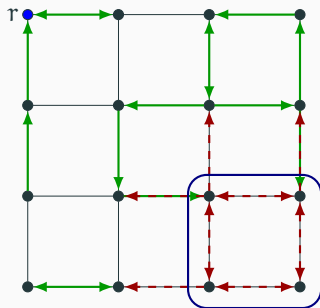


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Resampling table

Associate an infinite stack $X_{i,0}, X_{i,1}, \dots$ to each random variable X_i . When we need to resample, draw the next value in the stack.

X_1	$X_{1,0}$	$X_{1,1}$	$X_{1,2}$	$X_{1,3}$	$X_{1,4}$	\dots
X_2	$X_{2,0}$	$X_{2,1}$	$X_{2,2}$	$X_{2,3}$	$X_{2,4}$	\dots
X_3	$X_{3,0}$	$X_{3,1}$	$X_{3,2}$	$X_{3,3}$	$X_{3,4}$	\dots
X_4	$X_{4,0}$	$X_{4,1}$	$X_{4,2}$	$X_{4,3}$	$X_{4,4}$	\dots

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Change the future, not the history

For **extremal** instances, replacing a **perfect** assignment with another one will not change the resampling history!

X_1	$X_{1,0}$	$X_{1,1}$	$X_{1,2}$	$X_{1,3}$	$X_{1,4}$	\dots
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X_3		Λ_2		$X_{3,3}$	$X_{3,4}$	\dots
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X_1	$X'_{1,0}$	$X_{1,1}$	$X_{1,2}$	$X_{1,3}$	$X_{1,4}$	\dots
X_2	A_1	$X'_{2,1}$	$X_{2,2}$	$X_{2,3}$	$X_{2,4}$	\dots
X_3		A_2	$X'_{3,2}$	$X_{3,3}$	$X_{3,4}$	\dots
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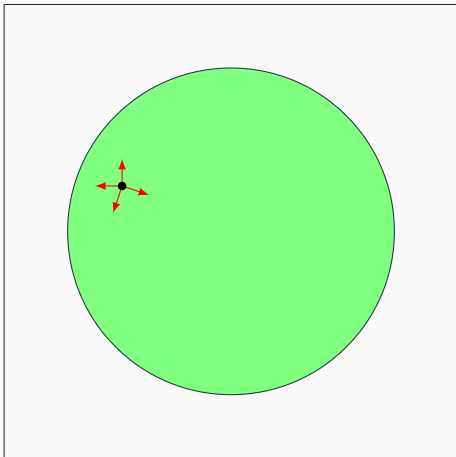
X_1	$X'_{1,0}$	$X_{1,1}$	$X_{1,2}$	$X_{1,3}$	$X_{1,4}$	\dots
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For any two outputs σ and τ , there is a **bijection** between trajectories leading to σ and τ .

Partial Rejection Sampling vs Markov chains

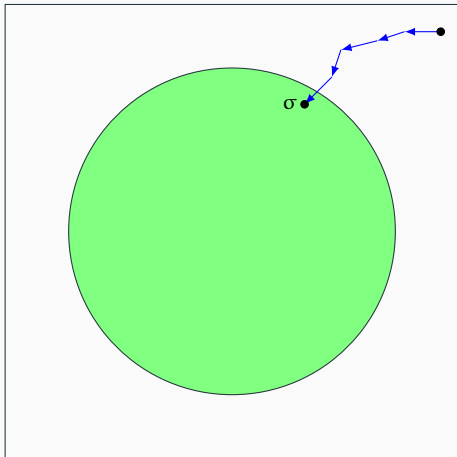
Markov chain is a random walk in the **solution** space.

(The solution space has to be connected,
and the mixing time is not easy to analyze.)



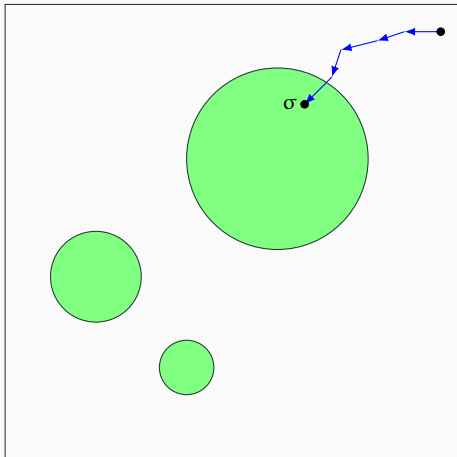
Partial Rejection Sampling vs Markov chains

PRS is a local search on the **whole** space.



Partial Rejection Sampling vs Markov chains

PRS is a local search on the **whole** space.
(Ergodicity is not an issue.)

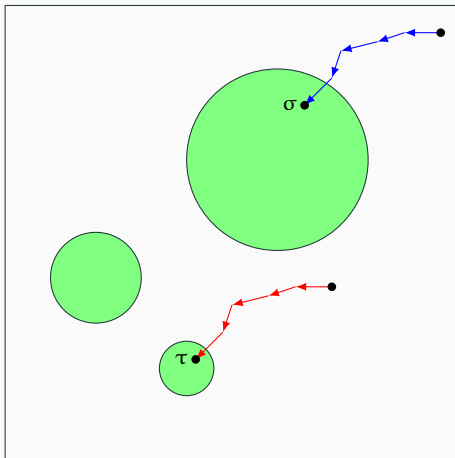


Partial Rejection Sampling vs Markov chains

PRS is a local search on the **whole** space.

(Correctness guaranteed by the bijection.)

Exact formula for its running time on extremal instances.)



Theorem (G., Jerrum, and Liu, 2017)

Under Shearer's condition, for *extremal* instances,

$$\mathbb{E} T = \frac{\text{total weight of one-flaw assignments}}{\text{total weight of perfect assignments}}.$$

(Shearer (1985) has shown a sufficient condition to guarantee the existence of one perfect assignment, which is optimal for Lovász Local Lemma.)

The upper bound is shown by Kolipaka and Szegedy (2011).

Back to cluster-popping

Cluster-popping: repeatedly resample minimal clusters.

Let Ω_k be the set of subgraphs with k minimal clusters, and

$$Z_k := \sum_{S \in \Omega_k} p^{|\mathbb{E} \setminus S|} (1-p)^{|S|}. \quad \text{Then, } \mathbb{E} T = \frac{Z_1}{Z_0}.$$

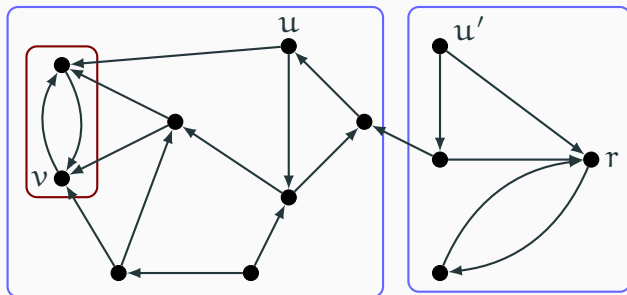
Lemma (G. and Jerrum, 2018)

For *bi-directed* graphs, $Z_1 \leq \frac{p}{1-p} \cdot mnZ_0$.

We show this by designing an injective mapping $\Omega_1 \rightarrow \Omega_0 \times V \times E$.

Injective mapping

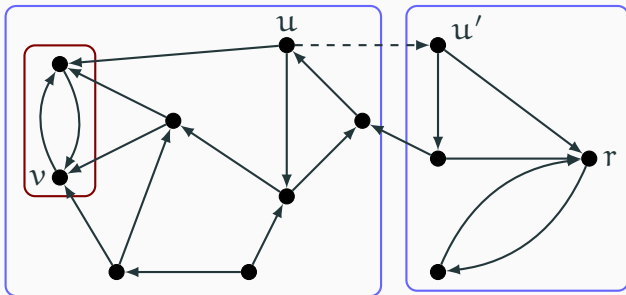
Given $R \in \Omega_1$, we map it to $R_0 \in \Omega_0$ by “repairing” the unique minimal cluster.



Conversely, given $R_0 \in \Omega_0$, (u, u') and v , we can recover $R \in \Omega_1$.

Injective mapping

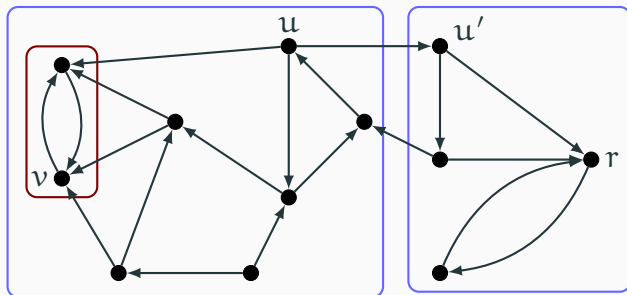
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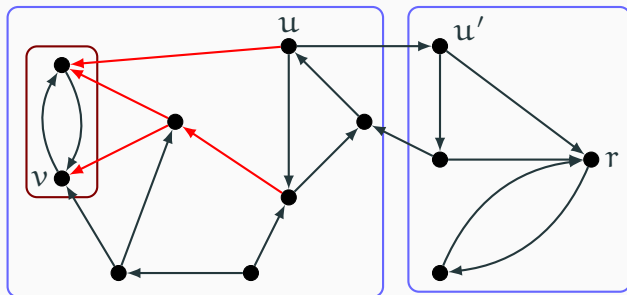
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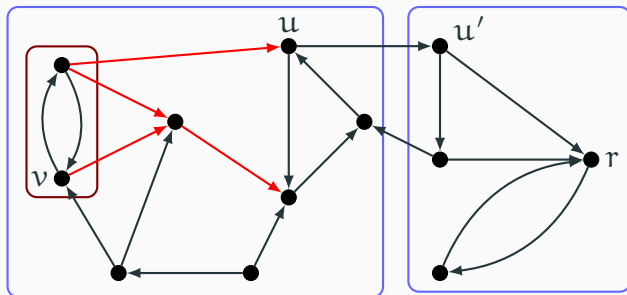
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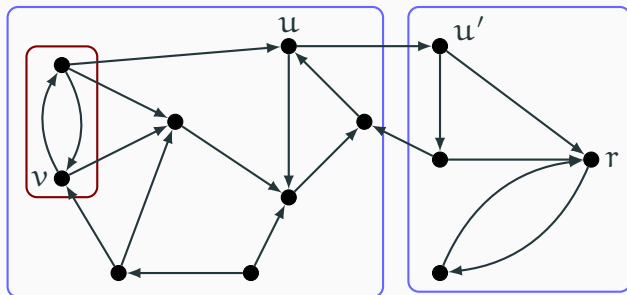
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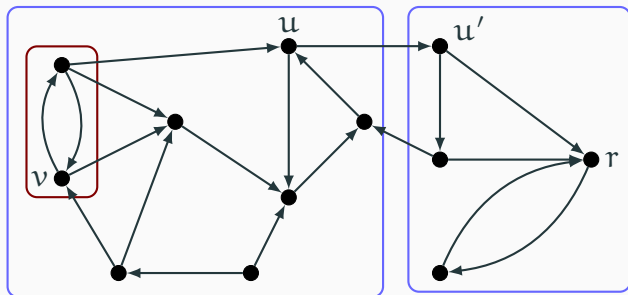
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Injective mapping

Given $R \in \Omega_1$, we map it to $R_0 \in \Omega_0$ by “repairing” the unique minimal cluster.



Conversely, given $R_0 \in \Omega_0$, (u, u') and v , we can recover $R \in \Omega_1$.

Recap for reliability

Approximate $Z_{\text{rel}}(G)$ via a sequence of contractions G_0, \dots, G_{n-1} , and estimate each $\frac{Z_{\text{rel}}(G_i)}{Z_{\text{rel}}(G_{i+1})}$ using the following sampling oracle:

1. run cluster-popping to sample a root-connected subgraph in \vec{G} ;
2. use the coupling to get a random connected subgraph.

To bound the running time of cluster-popping, we use a result of (GJL'17) and design an injective mapping.

Counting connected subgraphs of fixed size

Let S_t be the set of connected subgraph of size t where $n - 1 \leq t \leq m$ and $N_t = |S_t|$. Then a result of [Huh and Katz \(2012\)](#) implies that the sequence $(N_t)_t$ is log-concave, namely,

$$N_{t-1}N_{t+1} \leq N_t^2.$$

(The complements of connected subgraphs are independent sets of the co-graphic matroid associated with G , and co-graphic matroids are representable. So [HK'12](#) applies. Similar log-concavity in general matroids is resolved by [Adiprasito, Huh, and Katz \(2015\)](#).)

Given the sampler for connected subgraphs and log-concavity, we can set $p = \frac{N_t}{N_{t-1} + N_t}$ so that subgraphs in S_t show up frequently enough. There is a standard approach ([Jerrum and Sinclair, 1989](#)) to estimate each individual N_t .

Other examples of PRS

Extremal instances:

- Uniform spanning trees — cycle-popping ([Wilson, 1996](#))
- Uniform sink-free orientations — sink-popping ([Bubley and Dyer, 1997](#)) ([Cohn, Pemantle, and Propp, 2002](#))
- Uniform bases of bicircular matroids ([G. and Jerrum, 2018+](#))

General instances ([G., Jerrum, and Liu, 2017](#)):

- Weighted independent set (Hardcore gas model)
- Hard disks / hard spheres model ([G. and Jerrum, 2018](#))
- Solutions to k -CNF formulas with bounded variable degrees

Results for general instances are far from optimal.

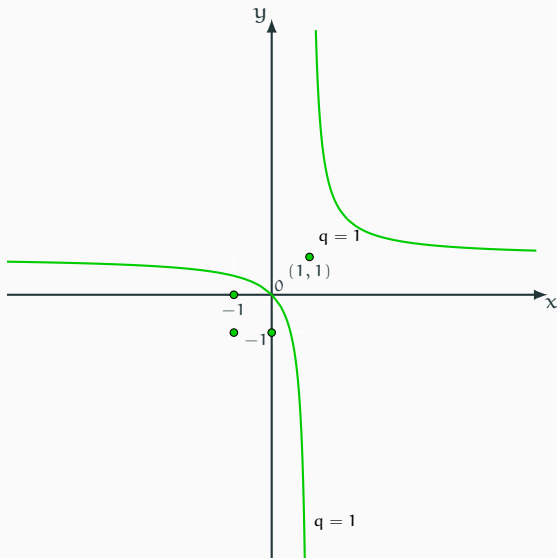
Can we do this for colourings?

CONCLUDING REMARKS

Approximating the Tutte polynomial

$$q = (x - 1)(y - 1)$$

Poly-time



Ref:

Jaeger, Vertigan, and Welsh (1990);

Jerrum and Sinclair (1993);

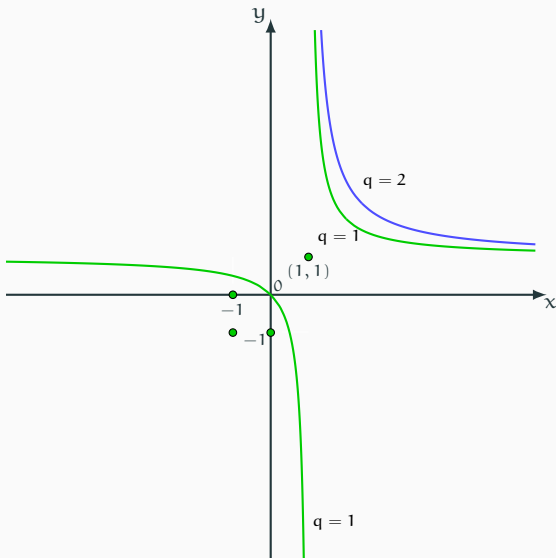
Goldberg and Jerrum (2008, 2012, 2014)

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FPRAS



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NP-hard to approximate
(#P-hard mostly)

#PM-equivalent

#BIS-hard

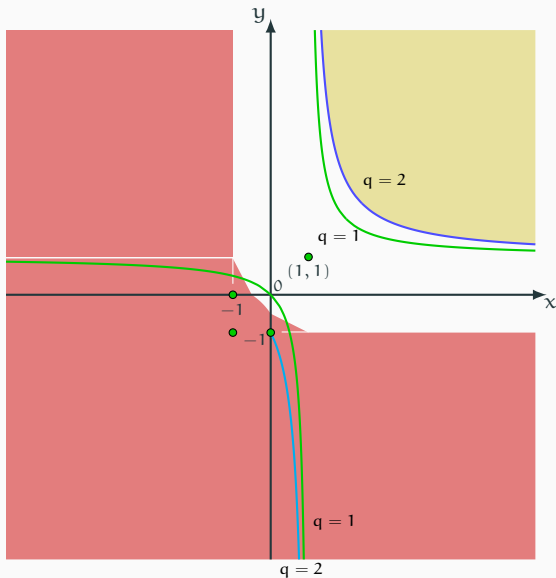
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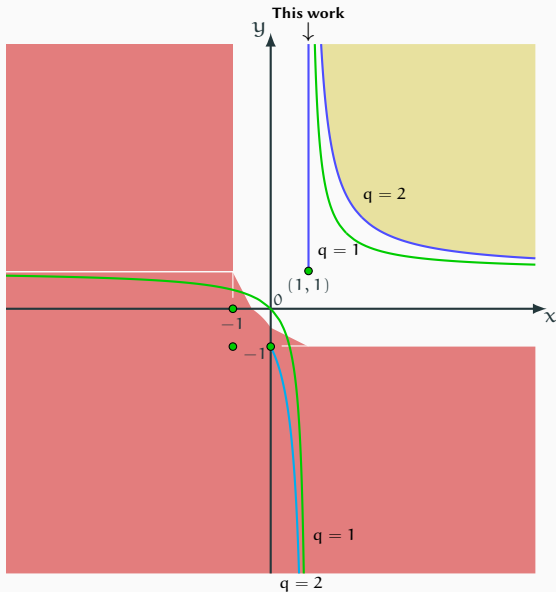
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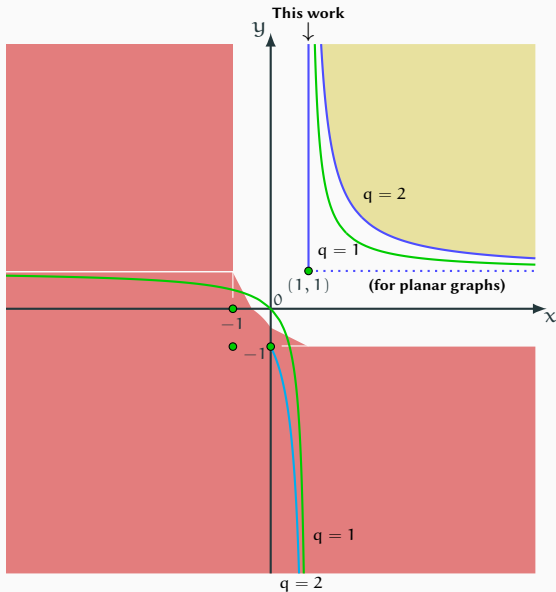
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A common paradigm

Both our result and the previous positive result on the Tutte plane (Jerrum and Sinclair, 1993) follow the same pattern:

1. Transform the problem into an equivalent one:
 - Ferromagnetic Ising model \rightarrow even subgraphs (JS'93);
 - Reliability \rightarrow bi-directed reachability.
2. Exploit some nice properties of the new solution space.

Are there other equivalences we have not discovered yet?

Open problems

- Is the **Markov chain** for connected subgraphs rapidly mixing?
- Approximating **s-t RELIABILITY**, and other variants?
(The natural Markov chain is exponentially slow for s-t version.)
- Approximating $Z_{\text{Tutte}}(\mathbf{G}; \chi, 1)$ for $\chi > 1$ (edge-weighted forests)?

THANK YOU!

arXiv:1611.01647

(PARTIAL REJECTION SAMPLING)

arXiv:1709.08561

(NETWORK RELIABILITY)

arXiv:1807.01680

(TIGHT BOUNDS)