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

Heng Guo (University of Edinburgh)

Joint with Mark Jerrum (Queen Mary, University of London)

Zhejiang University, Jun 07, 2018

P: polynomial-time computable NP: polynomial-time verifiable

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

a counting analogue of **NP**.

Evaluation of probabilities; Partition functions in statistical mechanics; Counting discrete structures ...



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

- at most NP-hard (Stockmeyer, 1985);
- any polynomial approximation can be amplified into an $(1 \pm \epsilon)$ -approximation with overhead polynomial in $1/\epsilon$.

Efficient approximation algorithms do exist! Most famous example: the permanent of a non-negative matrix (Jerrum, Sinclair, and Vigoda, 2004).

Many problems were proposed in 80s, and subsequently solved in 90s, but there are still a fair amount of leftovers!

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.

One may be curious about all kinds of properties of G(p):

- Pr[G(p) is connected] (ALL-TERMINAL) RELIABILITY
- Pr[s and t are connected in G(p)] s-t RELIABILITY
- Pr[G(p) is acyclic] ...

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

In other words, we want to compute

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

For example:

$$\begin{aligned} \mathsf{Z}_{rel}(&=, p) &= = = (1-p)^{n-1};\\ \mathsf{Z}_{rel}(&, p) &= &+ &+ &+ &+ &+ &\\ &= (1-p)^4 + 4p(1-p)^3;\\ \mathsf{Z}_{rel}(G, 1/2) &= \frac{|\{\mathsf{R} \subseteq \mathsf{E} : (\mathsf{V}, \mathsf{R}) \text{ is connected}\}|}{2^{|\mathsf{E}|}} \end{aligned}$$

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),

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

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

A few specializations of (x, y):

- (1,1): # of spanning trees;
- (1,2): # of connected subgraphs, and

$$Z_{Tutte}(G; 1, 1/p) = Z_{rel}(G, p) \cdot \frac{p^{|V| - |E| - 1}}{(1 - p)^{|V| - 1}};$$

- (2,1): # of forests (acyclic subgraphs);
- (x-1)(y-1) = 2 and x, y > 0: ferromagnetic Ising model.

The Tutte polynomial

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

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

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

A few specializations of (x, y):

- (1,1): # of spanning trees;
- (1,2): # of connected subgraphs, and

$$Z_{Tutte}(G;1,1/p) = Z_{rel}(G,p) \cdot \frac{p^{|V|-|E|-1}}{(1-p)^{|V|-1}};$$

- (2,1): # of forests (acyclic subgraphs);
- (x-1)(y-1) = 2 and x, y > 0: ferromagnetic Ising model.

The Tutte polynomial

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

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

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

A few specializations of (x, y):

- (1,1): # of spanning trees;
- (1,2): # of connected subgraphs, and

 $Z_{Tutte}(G; 1, 1/p) = Z_{rel}(G, p) \cdot \frac{p^{|V| - |E| - 1}}{(1 - p)^{|V| - 1}};$

- (2,1): # of forests (acyclic subgraphs);
- (x-1)(y-1) = 2 and x, y > 0: ferromagnetic Ising model.

Let $\mathfrak{m} := |\mathsf{E}|$ and $\mathfrak{n} := |\mathsf{V}|$.

Theorem (G. and Jerrum, 2018)

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

Theorem (G. and Jerrum, 2018)

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

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

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

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

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

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

When that happens, the variance of $l_{conn}(R)$ is exponentially large, and k has to be exponentially large to yield a good approximation.

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

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

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

It is easy to see that $\mathbb{E} \widetilde{Z} = \mathbb{Z}_{rel}$.

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

When that happens, the variance of $1_{conn}(R)$ is exponentially large, and k has to be exponentially large to yield a good approximation.

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

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

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

It is easy to see that $\mathbb{E} \widetilde{Z} = \mathbb{Z}_{rel}$.

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

When that happens, the variance of $1_{conn}(R)$ is exponentially large, and k has to be exponentially large to yield a good approximation.

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

Karger's algorithm has been subsequently refined by Harris and Srinivasan (2014), Karger (2016, 2017).

Karger (2017) is a recursive algorithm using NMC running in $O(n^{2.87})$.

- Run NMC, if $p^c > 1/2$, where c is the size of the min-cut.
- Otherwise, draw subgraphs $H_1, H_2 \sim G(q)$ where $q = 2^{-1/c} > p$, and $\frac{1}{2} (Z_{unrel}(H_1, p/q) + Z_{unrel}(H_2, p/q))$

is an unbiased estimator of $Z_{unrel}(G,p)$.

• Recursively estimate $Z_{unrel}(H_i, p/q)$ for i = 1, 2.

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

Karger's algorithm has been subsequently refined by Harris and Srinivasan (2014), Karger (2016, 2017).

Karger (2017) is a recursive algorithm using NMC running in $O(n^{2.87})$.

- Run NMC, if $p^c > 1/2$, where c is the size of the min-cut.
- Otherwise, draw subgraphs $H_1, H_2 \sim G(q)$ where $q = 2^{-1/c} > p$, and $\frac{1}{2} (Z_{unrel}(H_1, p/q) + Z_{unrel}(H_2, p/q))$

is an unbiased estimator of $Z_{unrel}(G, p)$.

• Recursively estimate $Z_{unrel}(H_i, p/q)$ for i = 1, 2.

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

Karger's algorithm has been subsequently refined by Harris and Srinivasan (2014), Karger (2016, 2017).

Karger (2017) is a recursive algorithm using NMC running in $O(n^{2.87})$.

- Run NMC, if $p^c > 1/2$, where c is the size of the min-cut.
- Otherwise, draw subgraphs $H_1, H_2 \sim G(q)$ where $q = 2^{-1/c} > p$, and $\frac{1}{2} (Z_{unrel}(H_1, p/q) + Z_{unrel}(H_2, p/q))$

is an unbiased estimator of $Z_{unrel}(G,p)$.

• Recursively estimate $Z_{unrel}(H_i, p/q)$ for i = 1, 2.

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

Karger's algorithm has been subsequently refined by Harris and Srinivasan (2014), Karger (2016, 2017).

Karger (2017) is a recursive algorithm using NMC running in $O(n^{2.87})$.

- Run NMC, if $p^c > 1/2$, where c is the size of the min-cut.
- Otherwise, draw subgraphs $H_1, H_2 \sim G(q)$ where $q=2^{-1/c}>p$, and $\frac{1}{2}\left(Z_{unrel}(H_1,p/q)+Z_{unrel}(H_2,p/q)\right)$

is an unbiased estimator of $Z_{unrel}(G, p)$.

- Recursively estimate $Z_{\text{unrel}}(H_i,p/q)$ for i=1,2.

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

Karger's algorithm has been subsequently refined by Harris and Srinivasan (2014), Karger (2016, 2017).

Karger (2017) is a recursive algorithm using NMC running in $O(n^{2.87})$.

- Run NMC, if $p^c > 1/2$, where c is the size of the min-cut.
- Otherwise, draw subgraphs $H_1, H_2 \sim G(q)$ where $q=2^{-1/c}>p$, and $\frac{1}{2}\left(Z_{unrel}(H_1,p/q)+Z_{unrel}(H_2,p/q)\right)$

is an unbiased estimator of $Z_{unrel}(G, p)$.

- Recursively estimate $Z_{\text{unrel}}(H_i,p/q)$ for i=1,2.

Let $\pi_G(\cdot)$ be the product distribution over the edges, conditioned on the resulting graph being connected.

Let $\pi_G(\cdot)$ be the product distribution over the edges, conditioned on the resulting graph being connected.



Let $\pi_G(\cdot)$ be the product distribution over the edges, conditioned on the resulting graph being connected.



Let $\pi_G(\cdot)$ be the product distribution over the edges, conditioned on the resulting graph being connected.



Let $\pi_G(\cdot)$ be the product distribution over the edges, conditioned on the resulting graph being connected.



Let $\pi_G(\cdot)$ be the product distribution over the edges, conditioned on the resulting graph being connected.

We can approximate Z_{rel} using an oracle drawing from π_G .



Rewrite

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

Let $\pi_G(\cdot)$ be the product distribution over the edges, conditioned on the resulting graph being connected.

We can approximate Z_{rel} using an oracle drawing from π_G .



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

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

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

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

- **1.** Let $C_0 = E$.
- $\label{eq:constraint} \mbox{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, nothing 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.



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

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

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

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

where \overrightarrow{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 C under which

G(p) is connected $\Leftrightarrow \overrightarrow{G}(p)$ 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.

A coupling proof

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

There is a coupling C under which

```
G(p) is connected \Leftrightarrow \overrightarrow{G}(p) is root-connected.
```

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



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

A coupling proof

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

There is a coupling C under which

```
G(p) is connected \Leftrightarrow \overrightarrow{G}(p) 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 \mathbf{r} are exactly the same.

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, \ldots, 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.

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.)

- 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, \ldots, 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.
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, \ldots, 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.

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, \ldots, 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.

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, \ldots, 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.










































































PARTIAL REJECTION SAMPLING

(A GENERAL THEORY BEHIND CLUSTER-POPPING)

Cluster-popping falls into the 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).

Cluster-popping under partial rejection sampling:

Arcs are variables. Minimal clusters are "bad" events.



There can be exponentially many bad events.

if any two "bad" events A_i and A_j are either independent or disjoint.



if any two "bad" events A_i and A_j are either independent or disjoint.



if any two "bad" events A_i and A_j are either independent or disjoint.



if any two "bad" events A_i and A_j are either independent or disjoint.



if any two "bad" events A_i and A_j are either independent or disjoint.



X ₁	X _{1,0}	X _{1,1}	X _{1,2}	X _{1,3}	X _{1,4}	
X ₂	X _{2,0}	X _{2,1}	X _{2,2}	X _{2,3}	X _{2,4}	•••
X ₃	X _{3,0}	X _{3,1}	X _{3,2}	X _{3,3}	X _{3,4}	•••
X_4	X _{4,0}	X _{4,1}	X _{4,2}	X _{4,3}	X _{4,4}	•••

X ₁	X _{1,0}	X _{1,1}	X _{1,2}	X _{1,3}	X _{1,4}	•••
X ₂	X _{2,0}	X _{2,1}	X _{2,2}	X _{2,3}	X _{2,4}	•••
X ₃	X _{3,0}	X _{3,1}	X _{3,2}	X _{3,3}	X _{3,4}	•••
X ₄	X _{4,0}	X _{4,1}	X _{4,2}	X _{4,3}	X _{4,4}	•••

X ₁	X _{1,0}	X _{1,1}	X _{1,2}	X _{1,3}	X _{1,4}	•••
X ₂	X _{2,0}	X _{2,1}	X _{2,2}	X _{2,3}	X _{2,4}	•••
X ₃	X _{3,0}	X _{3,1}	X _{3,2}	X _{3,3}	X _{3,4}	•••
X ₄	X _{4,0}	X _{4,1}	X _{4,2}	X _{4,3}	X _{4,4}	•••

X ₁	X _{1,0}	X _{1,1}	X _{1,2}	X _{1,3}	X _{1,4}	•••
X ₂		X _{2,1}	X _{2,2}	X _{2,3}	X _{2,4}	•••
X ₃		X _{3,1}	X _{3,2}	X _{3,3}	X _{3,4}	•••
X ₄	X _{4,0}	X _{4,1}	X _{4,2}	X _{4,3}	X _{4,4}	•••

X ₁	X _{1,0}	X _{1,1}	X _{1,2}	X _{1,3}	X _{1,4}	•••
X ₂		X _{2,1}	X _{2,2}	X _{2,3}	X _{2,4}	•••
X ₃		X _{3,1}	X _{3,2}	X _{3,3}	X _{3,4}	•••
X ₄	X _{4,0}	X _{4,1}	X _{4,2}	X _{4,3}	X _{4,4}	•••

X ₁	X _{1,0}	X _{1,1}	X _{1,2}	X _{1,3}	X _{1,4}	•••
X ₂		X _{2,1}	X _{2,2}	X _{2,3}	X _{2,4}	•••
X ₃			X _{3,2}	X _{3,3}	X _{3,4}	•••
X ₄		X _{4,1}	X _{4,2}	X _{4,3}	X _{4,4}	•••

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}	•••
X ₂		X _{2,1}	X _{2,2}	X _{2,3}	X _{2,4}	•••
X ₃			X _{3,2}	X _{3,3}	X _{3,4}	•••
X ₄		X _{4,1}	X _{4,2}	X _{4,3}	X _{4,4}	•••

For extremal instances, replacing a perfect assignment with another one will not change the resampling 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}	•••
X ₂		X' _{2,1}	X _{2,2}	X _{2,3}	X _{2,4}	•••
X ₃		A_2	X' _{3,2}	X _{3,3}	X _{3,4}	•••
X ₄		X' _{4,1}	X _{4,2}	X _{4,3}	X _{4,4}	•••

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}	
X ₂		X' _{2,1}	X _{2,2}	X _{2,3}	X _{2,4}	
X ₃	A_1	A_2	X' _{3,2}	X _{3,3}	X _{3,4}	
X_4		X' _{4,1}	X _{4,2}	X _{4,3}	X _{4,4}	

For any output σ and τ , there is a bijection between trajectories leading to σ and τ .

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.)



PRS is a local search on the whole space.



PRS is a local search on the whole space.

(Ergodicity is not an issue.)



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).

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

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

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 $\Omega_1 \rightarrow \Omega_0 \times V \times E$.







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









Approximate $Z_{rel}(G)$ via a sequence of contractions G_0, \ldots, G_{n-1} , and estimate each $\frac{Z_{rel}(G_i)}{Z_{rel}(G_{i+1})}$ 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.
Let S_t be the set of connected subgraph of size t where $n-1\leqslant t\leqslant m$ and N_t = $|S_t|$. Then a result of Huh and Katz (2012) implies that the sequence $(N_t)_t$ is log-concave, namely,

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

(Here we treat connected subgraphs as independent sets in the dual of the graphic matroid, which is representable and HK'12 applies. Similar log-concavity in general matroid 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 .

Extremal instances:

- Uniform spanning trees cycle-popping (Wilson, 1996)
- Uniform sink-free orientations sink-popping (Bubley and Dyer, 1997) (Cohn, Pemantle, and Propp, 2002)

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.

CONCLUDING REMARKS

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

Poly-time

Ref: Jaeger, Vertigan, and Welsh (1990); Jerrum and Sinclair (1993); Goldberg and Jerrum (2008, 2012, 2014)





Poly-time

FPRAS





Approximating the Tutte polynomial

 $\boldsymbol{q}=(\boldsymbol{x}-1)(\boldsymbol{y}-1)$

Poly-time FPRAS NP-hard to approximate (#P-hard mostly)

#PM-equivalent

#BIS-hard

Open: white area

Ref: Jaeger, Vertigan, and Welsh (1990); Jerrum and Sinclair (1993); Goldberg and Jerrum (2008, 2012, 2014)



Approximating the Tutte polynomial

 $\boldsymbol{q}=(\boldsymbol{x}-1)(\boldsymbol{y}-1)$

Poly-time FPRAS NP-hard to approximate (#P-hard mostly)

#PM-equivalent

#BIS-hard

Open: white area

Ref:

Jaeger, Vertigan, and Welsh (1990); Jerrum and Sinclair (1993); Goldberg and Jerrum (2008, 2012, 2014)



Approximating the Tutte polynomial

 $\boldsymbol{q}=(\boldsymbol{x}-1)(\boldsymbol{y}-1)$

Poly-time FPRAS NP-hard to approximate (#P-hard mostly)

#PM-equivalent

#BIS-hard

Open: white area

Ref:

Jaeger, Vertigan, and Welsh (1990); Jerrum and Sinclair (1993); Goldberg and Jerrum (2008, 2012, 2014)



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?

- 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 $Z_{Tutte}(G; x, 1)$ for x > 1 (edge-weighted forests)?

THANK YOU!

arXiv:1611.01647 (Partial rejection sampling)

arXiv:1709.08561 (Network reliability)