

MARKOV CHAIN ALGORITHMS FOR BOUNDED DEGREE k -SAT

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One of the most important problem in computer science

Input: A formula in **conjunctive normal form**, like

$$(x_1 \vee \bar{x}_3 \vee x_5) \wedge (x_2 \vee x_3) \wedge (\bar{x}_3 \vee \bar{x}_4) \wedge (x_1 \vee \bar{x}_5 \vee x_6 \vee x_7) \dots$$

Output: Is it satisfiable?

The first **NP**-complete problem **Cook (1971)** – **Levin (1973)**

Sometimes we are not satisfied with finding one solution. We want to generate a **uniformly at random** solution.

The ability of sampling solutions enables us to

- approximately count the number of solutions;
- estimate the marginal probability of individual variables;
- estimate other quantities of interest ...

And sometimes generating random instances satisfying given constraints can be useful too.

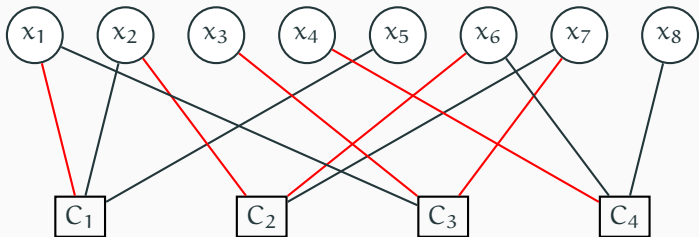
Sampling can be **NP**-hard even if finding a solution is easy (e.g. under Lovász local lemma conditions).

A NATURAL (BUT NOT WORKING) APPROACH

Standard sampling approach: Glauber dynamics / Gibbs sampling

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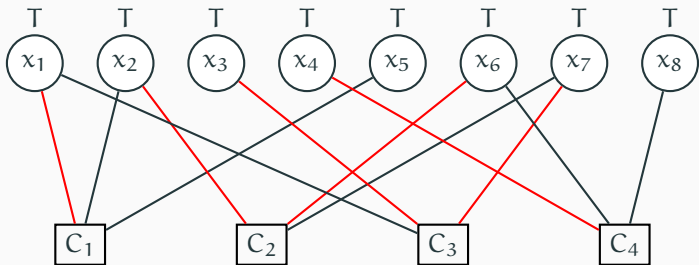


$$(\neg x_1 \vee x_2 \vee x_5) \wedge (\neg x_2 \vee \neg x_6 \vee x_7) \wedge (x_1 \vee \neg x_3 \vee \neg x_7) \wedge (\neg x_4 \vee x_6 \vee x_8)$$

Choose a random variable, sample its value conditioned on all others.

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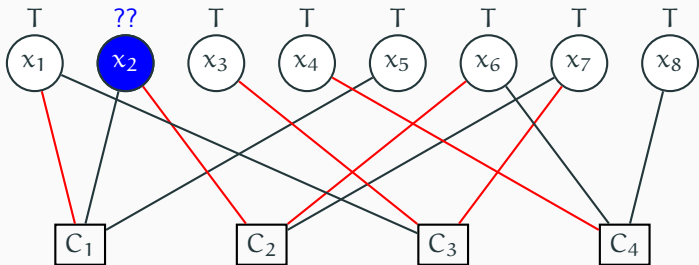


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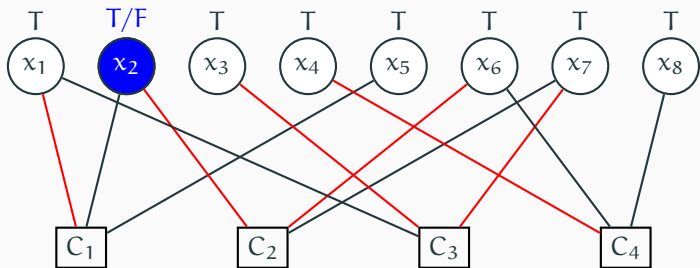


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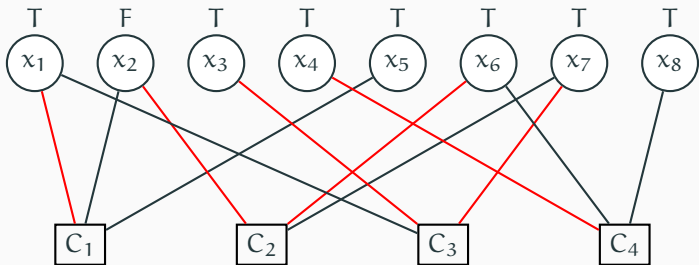


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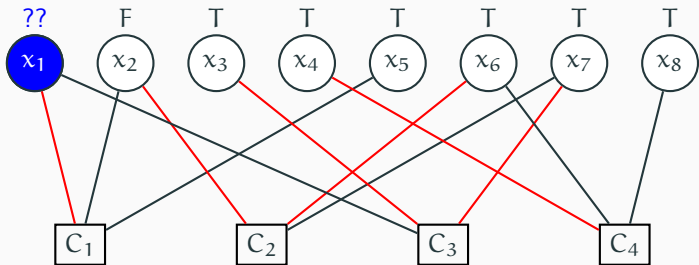


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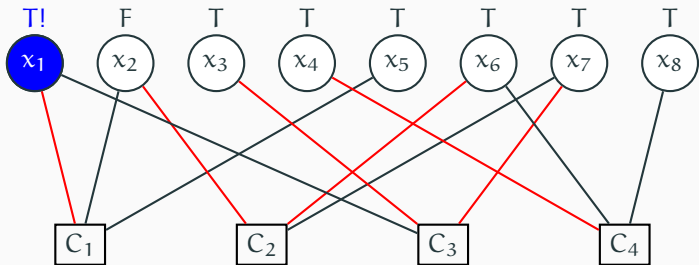


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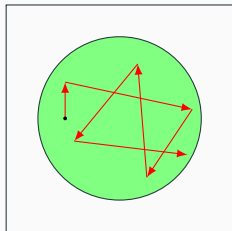
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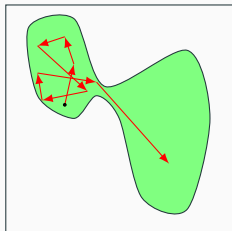
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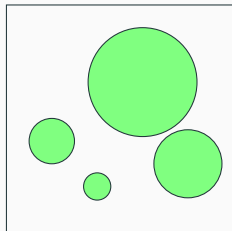
THREE SCENARIOS FOR MARKOV CHAINS



Fast mixing

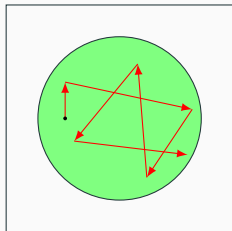


Slow mixing

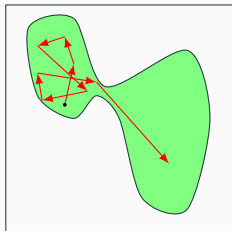


Not mixing!

THREE SCENARIOS FOR MARKOV CHAINS



Fast mixing



Slow mixing



Not mixing!

DISCONNECTIVITY FOR k-SAT

Suppose we have k variables, and each clause contains all k variables.

$$\Phi = C_1 \wedge C_2 \wedge \dots \wedge C_m$$

Each C_i forbids one assignment on k variables.

For example, $C_i = x_1 \vee x_2 \vee \dots \vee x_k$ forbids the all False assignment.

Thus, if we forbade all assignments of Hamming weight i for some $1 \leq i \leq k - 1$ (using $\binom{k}{i}$ clauses), the solution space is not connected via single variable updates.

For example, to remove Hamming weight $k - 1$ assignments, we only need clauses

$$C_1 = \neg x_1 \vee x_2 \vee \dots \vee x_k$$

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\vdots

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OUR SOLUTION — PROJECTION

Projecting from a high dimension to a lower dimension may improve connectivity.

We will run Glauber dynamics on the projected distribution over a suitably “**marked**” variables.

The general problem is **NP**-hard, so we will focus on bounded degree cases.

BOUNDED DEGREE k -SAT



Theorem (Lovász local lemma)

Let E_1, \dots, E_m be a set of “bad” events, such that $\Pr[E_i] \leq p$ for all i . Moreover, each E_i is independent from all but at most Δ events. If $ep\Delta \leq 1$, then

$$\Pr \left[\bigwedge_{i=1}^m \overline{E_i} \right] > 0.$$

In the setting of k -SAT, each clause C_i defines a bad event E_i , which is the forbidden assignment of C_i , and $p = 2^{-k}$.

If every variable appears in at most d clauses, then $\Delta \leq kd$.

$$\begin{aligned} ep\Delta \leq 1 &\Leftrightarrow e2^{-k}kd \leq 1 \\ &\Leftrightarrow k \geq \log d + \log k + C \end{aligned}$$

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MOSER-TARDOS ALGORITHM

We consider k -CNF formula with variable degree at most d .

Theorem (Moser and Tardos, 2011)

If $k \geq \log d + \log k + C$, then we can always find a satisfying assignment in polynomial time.

The algorithm is extremely simple: assign variables u.a.r., then keep resample variables in violating clauses.

Unfortunately, sampling is substantially harder.

Theorem (Bezáková, Galanis, Goldberg, G. and Štefankovič, 2016)

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OPEN PROBLEM:

*IS THERE AN EFFICIENT ALGORITHM TO
SAMPLE SATISFYING ASSIGNMENTS OF k -SAT
GIVEN $k \gtrsim 2 \log d + C$?*

Hermon, Sly and Zhang (2016) Glauber dynamics mixes in $O(n \log n)$ time if $k \geq 2 \log d + C$ and there is **no negation** (monotone formula).

G., Jerrum and Liu (2016) “Partial rejection sampling” terminates in $O(n)$ time if $k \geq 2 \log d + C$ and there is **no small intersection**.

Moitra (2016) An “exotic” deterministic algorithm in $n^{O(k^2 d^2)}$ time if $k \geq 60(\log d + \log k) + 300$.

Theorem (Our result)

We give a Markov chain based algorithm in $\tilde{O}(n^{1+\delta} k^3 d^2)$ time if $k \geq 20(\log d + \log k) + \log \delta^{-1}$ where $\delta \leq 1/60$ is an arbitrary constant.

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OUR ALGORITHM



Goal: to sample from the uniform distribution μ over satisfying assignments

1. Mark a set M of variables;
2. Run Glauber dynamics on the projected distribution μ_M for $O(n \log n)$ steps. This yields an (approximate) sample $\sigma_M \sim \mu_M$;
3. Use rejection sampling to sample from $\sigma_{V \setminus M}$;
4. Output $\sigma_M \cup \sigma_{V \setminus M}$.

A set M of variables are marked so that:

1. for any clause C_i , $|C_i \cap M| \gtrsim 0.11k$;
2. for any clause C_i , $|C_i \setminus M| \gtrsim 0.51k$;

The existence of M is guaranteed by the local lemma, and M can be found by the [Moser-Tardos](#) algorithm in linear time.

TWO SIDES OF THE MARKING

If $|C_i \cap M|$ is large, then all components are small.

Lemma

For almost all $\sigma \in \{0, 1\}^M$, $V \setminus M$ scatters into connected components of size $O(\text{poly}(dk) \log n)$.

If $|C_i \setminus M|$ is large, then all variables are close to the uniform distribution.

Lemma

Conditioned on any assignment of M , for any $v \in V \setminus M$,

$$\left| \Pr_{\sigma \sim \mu_{V \setminus M}} [\sigma(v) = 1] - \frac{1}{2} \right| \leq \exp(-O(k)).$$

So the marking is to balance these two effects.

1. The Glauber dynamics on the marked variables is rapidly mixing;
2. The Glauber dynamics on the marked variables can be implemented efficiently;
3. The rejection sampling step in the end terminates quickly.

Item (1) is shown using the **path coupling** method.

Items (2) and (3) are shown together. In particular, the Glauber dynamics is implemented using **rejection sampling**.

IMPLEMENTING THE GLAUBER DYNAMICS

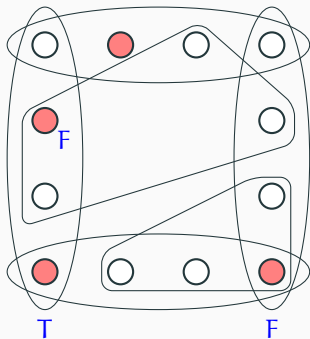
Glauber dynamics: compute the marginal probability of a variable conditioned on all other marked variables, which defines a smaller instance. (#P-hard in general.)

We approximately implement this by using rejection sampling on

1. all **unmarked** variables, and
2. the variable to be **updated**.

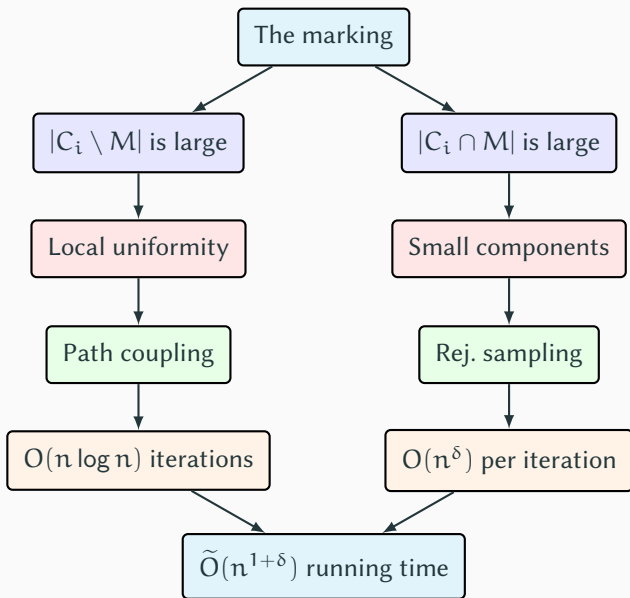
Rejection sampling terminates in $O(n^\delta)$ steps with high probability.

$$\Pr[\sigma(x_i) = T] = ??$$



(Every clause here has at least 1 marked variable and 2 unmarked variable.)

OVERVIEW



WHY REJECTION SAMPLING?

Draw $\sigma_M \sim \mu_M$. For each clause, there are at least $\Omega(k)$ variables assigned. Many clauses are satisfied, and the remaining clauses scatter into connected components of size $\asymp \log n$.

However, the size is $\Omega(dk \log n)$, so a brute-force enumeration takes time $n^{\Omega(dk)}$ which is too slow to our needs.

We use the local lemma once again here to show that uniform at random assignments satisfies remaining clauses with probability at least $\Omega(n^{-\delta})$. Thus, the rejection sampling succeeds in time $\tilde{O}(n^\delta)$.

PATH COUPLING

Path coupling condition: given two assignment σ_0 and σ_1 which differ on only one variable v ,

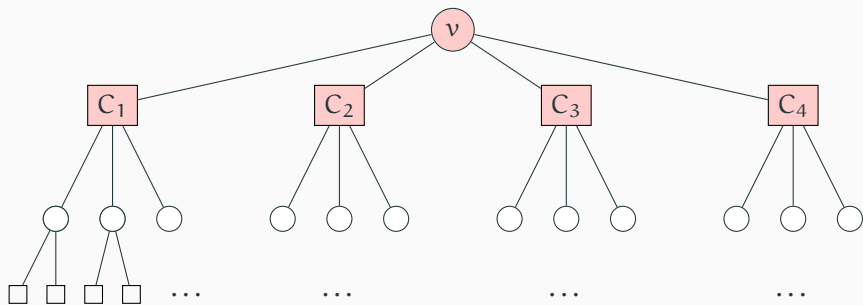
$$\sum_{u \in M, u \neq v} d_{TV}(\mu_u(\cdot \mid \sigma_0), \mu_u(\cdot \mid \sigma_1)) < 1.$$

Using the coupling inequality, it suffices to show that for a carefully designed coupling \mathcal{C} of $\mu_{M \setminus \{v\}}(\cdot \mid \sigma_0)$ and $\mu_{M \setminus \{v\}}(\cdot \mid \sigma_1)$ such that

$$\mathbb{E}_{(\tau_0, \tau_1) \sim \mathcal{C}} |\{\mathbf{u} \mid \mathbf{u} \in M \setminus \{v\}, \tau_0(\mathbf{u}) \neq \tau_1(\mathbf{u})\}| < 1.$$

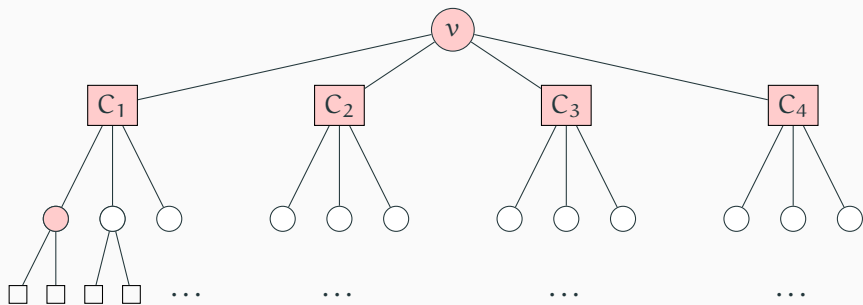
This “disagreement coupling” \mathcal{C} is very similar to the one used by [G., Liao, Lu and Zhang \(2018\)](#), which is a refined version of [Moitra \(2016\)](#). However, previous analysis has a **whp** guarantee, and we need a new analysis to bound the **expectation**.

DISAGREEMENT PERCOLATION



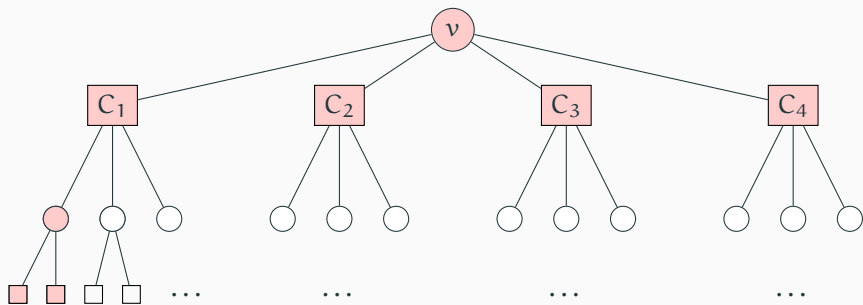
The disagreement percolation is similar to a branching process with branching factor dk , and each child survives with probability $\exp(-O(k))$ due to local uniformity.

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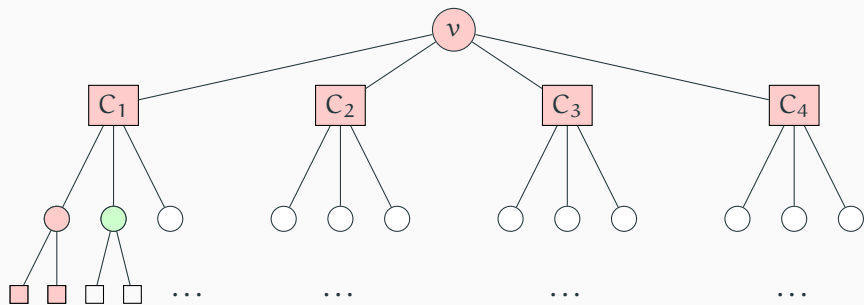
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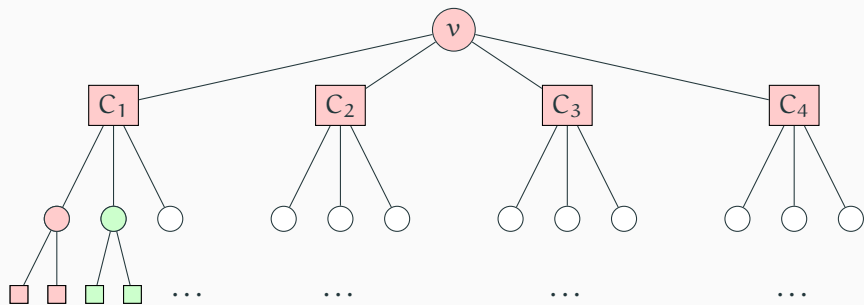
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CONCLUDING REMARKS



In another recent work, [Galanis, Goldberg, G. and Yang \(2019\)](#) showed that there is an efficient algorithm to approximately count the number of satisfying assignment of a **random** k -SAT instance with high probability, if the density is at most $2^{k/300}$.

- This improves the previous best algorithm which works for density $\leq \frac{2 \log k}{k}$ ([Montanari and Shah 2007](#)).
- The algorithm is based on [Moitra \(2016\)](#), with some extra ingredients to handle $\Omega(\log n)$ degree variables.
- Nonetheless, it is not clear if the Markov chain approach works for random formulas.

- Is the conjectured threshold correct?
 - Getting rid of the marking?
- Other CSPs, like hypergraph colouring?
- Other applications of this projection method?

Thank you!

arXiv:1911.01319